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970 10.6 50. 970 10.6 50. 971 10.6 50. 972 10.6 50. 973 10.6 50. 974 10.6 50. 975 10.6 50.	964 10.6 50. 965 10.6 50. 966 10.6 50. 967 10.6 50. 968 10.6 50.	959 10.8 51. 960 10.6 50. 961 10.6 50. 962 10.6 50.	955 10.8 51. 956 10.8 51. 957 10.8 51. 958 10.8 51.	952 10.8 51. 953 10.8 51. 954 10.8 51.	949 10.8 950 10.8 951 10.8	946 10.8 51. 947 10.8 51. 948 10.8 51.	944 10.8 51. 945 10.8 51.	941 10.8 51. 942 10.8 51.	938 10.8 51. 939 10.8 51.	935 10.8 936 10.8 937 10.8	10.8 51. 10.8 51. 10.8 51.	930 10.8 51. 931 10.8 51.	927 10.8 928 10.8 929 10.8	925 10.8 51. 926 10.8 51.	10.8 51.	10.8 51. 10.8 51. 10.8 51.	10.8 51. 10.8 51.	10.8 51. 10.8 51.	914 10.8 51. 915 10.8 51.	913 10.8 51.	911 10.8 51.	909 10.8 51. 910 10.8 51.	907 10.8 51. 908 10.8 51.	904 10.8 51. 905 10.8 51. 906 10.8 51.
969 10.6 50.5 20 970 10.6 50.5 20 971 10.6 50.5 20 972 10.6 50.5 20 973 10.6 50.5 20 974 10.6 50.5 21 975 10.6 50.5 21 976 10.6 50.5 21	964 10.6 50.5 20 964 10.6 50.5 20 965 10.6 50.5 20 966 10.6 50.5 20 967 10.6 50.5 20 968 10.6 50.5 20	959 10.8 51.4 70 960 10.6 50.5 17 961 10.6 50.5 17 962 10.6 50.5 18	955 10.8 51.4 68 956 10.8 51.4 68 957 10.8 51.4 70 958 10.8 51.4 70	952 10.8 51.4 68 953 10.8 51.4 68 954 10.8 51.4 68	949 10.8 51.4 60 950 10.8 51.4 60 951 10.8 51.4 68	946 10.8 51.4 53 947 10.8 51.4 53 948 10.8 51.4 56	944 10.8 51.4 53 945 10.8 51.4 53	941 10.8 51.4 48 942 10.8 51.4 50	938 10.8 51.4 47 939 10.8 51.4 47 940 10.8 51.4 47	935 10.8 51.4 47 936 10.8 51.4 47 937 10.8 51.4 47	10.8 51.4 45 10.8 51.4 47 10.8 51.4 47	930 10.8 51.4 43 931 10.8 51.4 44	927 10.8 51.4 42 928 10.8 51.4 43 929 10.8 51.4 43	925 10.8 51.4 40 926 10.8 51.4 41	10.8 51.4 40 10.8 51.4 40	10.8 51.4 40 10.8 51.4 40 10.8 51.4 40	10.8 51.4 40 10.8 51.4 40	10.8 51.4 39 10.8 51.4 39	914 10.8 51.4 36 915 10.8 51.4 36	913 10.8 51.4 36	911 10.8 51.4 35 912 10.8 51.4 36	909 10.8 51.4 35 910 10.8 51.4 35	907 10.8 51.4 35 908 10.8 51.4 35	904 10.8 51.4 35 905 10.8 51.4 35 906 10.8 51.4 35
969 10.6 50.5 970 10.6 50.5 971 10.6 50.5 972 10.6 50.5 973 10.6 50.5 974 10.6 50.5 975 10.6 50.5 975 10.6 50.5	964 10.6 50.5 20 3 965 10.6 50.5 20 3 966 10.6 50.5 20 3 967 10.6 50.5 20 4 968 10.6 50.5 20 4	959 10.8 51.4 70 3 960 10.6 50.5 17 1 961 10.6 50.5 17 1 962 10.6 50.5 18 4	955 10.8 51.4 68 3 956 10.8 51.4 70 1 957 10.8 51.4 70 1 958 10.8 51.4 70 2	952 10.8 51.4 68 1 953 10.8 51.4 68 1 954 10.8 51.4 68 2	949 10.8 51.4 60 1 950 10.8 51.4 60 2 951 10.8 51.4 68 1	946 10.8 51.4 53 4 947 10.8 51.4 53 5 948 10.8 51.4 56 1	944 10.8 51.4 53 3 945 10.8 51.4 53 3	941 10.8 51.4 48 1 942 10.8 51.4 50 4	938 10.8 51.4 47 4 939 10.8 51.4 47 4 940 10.8 51.4 47 4	935 10.8 51.4 47 4 936 10.8 51.4 47 4 937 10.8 51.4 47 4	10.8 51.4 45 1 10.8 51.4 47 3 10.8 51.4 47 4	930 10.8 51.4 43 3 931 10.8 51.4 44 1	927 10.8 51.4 42 1 928 10.8 51.4 43 2 929 10.8 51 4 43 2	925 10.8 51.4 40 4 926 10.8 51.4 41 2	10.8 51:4 40 3 10.8 51.4 40 3	10.8 51.4 40 3 10.8 51.4 40 3 10.8 51.4 40 3	10.8 51.4 40 2 10.8 51.4 40 3	10.8 51.4 39 1 10.8 51.4 39 1	915 10.8 51.4 36 4 915 10.8 51.4 36 4	913 10.8 51.4 36 4	911 10.8 51.4 35 6 912 10.8 51.4 36 3	909 10.8 51.4 35 5 910 10.8 51.4 35 5	907 10.8 51.4 35 4 908 10.8 51.4 35 5	904 10.8 51.4 35 1 905 10.8 51.4 35 1 906 10.8 51.4 35 3

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APPLICANT: Groner, Bernd
APPLICANT: Gouilleux, Fab:
APPLICANT: Wakao, Hiroshi
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                                                                 REGISTRATION NUMBER: 4-19
REFERENCE/DOCKET NUMBER: 4-19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEPAX: (919) 541-8689
TELEFAX: (919) 541-8689
                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: GB 940
FILING DATE: 11-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Elmer, James Scoot
REGISTRATION NUMBER: 36,12
                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
COMPUTER: IBM PC com
                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Cytokine Regulated Transcription Factor NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
           MOLECULE TYPE:
                  TYPE: nucleic STRANDEDNESS: TOPOLOGY: lin
                                                                                                                                                                                                                                                                       COUNTRY:
ZIP: 105
                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                      ADDRESSEE: CLDA Drive
                                                                                                                                                                                           FILING DATE
                                                                                                                                                                                                    APPLICATION NUMBER:
 DESCRIPTION:
                                                 LENGTH:
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                                       nucleic acid
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                                                  21 base pairs
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                   linear
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/desc = "MGF binding sequence from
                                                                                                                                                                                                                                                    Floppy disk
                               single
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US-08-148-117A-4
US-08-633-743-24
US-08-943-731-522
US-08-943-941-21
US-08-891-848-10
US-08-891-848-10
US-08-891-86A-8
PCT US92-01015-5
5231168-16-592-704
US-08-117-952-704
US-08-117-952-704
US-08-122-13
US-08-454-057-44
US-08-32-818-8
US-08-454-057-17
US-08-244-122-13
US-08-244-122-13
US-08-38-215-15-15
US-08-38-15-15-15
US-08-38-15-15-15
US-08-38-15-15-15
US-08-185-336-17
US-08-915-336-17
US-08-915-336-17
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US-08-915-336-17
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10, Appl
11, Appl
3, Appl
3, Appl
5, Appl
5, Appl
6, Appl
6, Appl
11, Appl
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24, Appl
522, App
55, Appl
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GENERAL INFORMATION:
APPLICANT: No. 6307024ak, Julia E.
APPLICANT: Presnell, Scott R.
APPLICANT: Sprecher, Cindy A.
APPLICANT: Foster, Donald C.
APPLICANT: Foster, Donald C.
APPLICANT: Gross, Jane A.
APPLICANT: Johnston, Janet V.
APPLICANT: Johnston, Janet V.
APPLICANT: Helson, Andrew J.
APPLICANT: Hammond, Angela K.
TITLE OF INVENTION: NOVEL CYTOKINE ZALPHA11 LIGAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; DESCRIPTION:
; HYPOTHETICAL:
US-08-366-276-8
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APPLICANT: No. 6307024ak, Julia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local Similarity
                                                                                                                                                                                                                   Sequence 60, Application US/09522217 Patent No. 6307024
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2000-03-09
EARLIER APPLICATION NUMBER: US 60/123,547
EARLIER FILING DATE: 1999-03-09
EARLIER FILING DATE: 1999-03-11
EARLIER FILING DATE: 1999-03-11
EARLIER APPLICATION NUMBER: US 60/142,013
EARLIER FILING DATE: 1999-07-01
NUMBER OF SEC ID NOS: 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/522,217
CURRENT FILING DATE: 2000-03-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Johnston, Janet V.
APPLICANT: Nelson, Andrew J.
APPLICANT: Dillon, Stacey R.
APPLICANT: Hammond, Angela K.
APPLICANT: Hammond, Angela K.
APPLICANT: Hammond, NOVEL CYTOKINE ZALPHAll LIGAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Presnell, Scott R.
APPLICANT: Sprecher, Cindy A.
APPLICANT: Foster, Donald C.
APPLICANT: Holly, Richard D.
APPLICANT: Gross, Jane A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 99-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 100
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
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6307024
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Pred. No.
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US-08-369-754-1
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US-08-369-754-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/522,217
CURRENT FILING DATE: 2000-03-09
EARLIER APPLICATION NUMBER: US 60/123,547
EARLIER FILING DATE: 1999-03-09
EARLIER APPLICATION NUMBER: US 60/123,904
EARLIER FILING DATE: 1999-03-11
EARLIER APPLICATION NUMBER: US 60/142,013
EARLIER FILING DATE: 1999-07-01
NUMBER OF SEQ ID NOS: 115
                    Query Match
Best Local Similarity
          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application Patent No. 6537553
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 60
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                                                                                                                                                     TELEFAX: 619-535-8949
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                                                                                                                                                                                                                           COMPUTEK REMARKATION OF THE PROPRY DISK

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/369,754
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: FUKUDA, MINORU
APPLICANT: SAWADA, RITSUKO
APPLICANT: TSUBOI, SHIGERU
TITLE OF INVENTION: E-SELECTIN-DEPENDENT CELL ADHESION
TITLE OF INVENTION: ANTAGONISTS
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                      REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: CAMPBELL AND FLORES
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                                                                                                                  LENGTH: 24 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
FILING DATE: 06-JAN
CLASSIFICATION: 435
                                                                                     TOPOLOGY:
                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 4370 LA
CITY: SAN DIEGO
                                                                                                                                                                                                        TELEPHONE:
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                                                                                                                                                                                      619-535-8949
        Conservative
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                                                                                       linear
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100.0%; Pr
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; Pred. No. 13;
   Score 15.4; DB 4;
Pred. No. 1.8e+02;
0; Mismatches 1;
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US-09-173-043-22
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US-09-140-804-46
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APPLICANT: Raymond, Christopher K.
TITLE OF INVENTION: PLASMID CONSTRUCTION BY HOMOLOGOUS
TITLE OF INVENTION: RECOMBINATION
FILE REFERENCE: 97-27
CURRENT APPLICATION NUMBER: US/09/173,043
                                                                                                                                                                                                               EARLIER APPLICATION NUMBER: US 60/062,061
EARLIER FILING DATE: 1997-10-16
NUMBER OF SEQ ID NOS: 34
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 22
LENGTH: 58
                                                                   Matches
                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22, Application US/09173043 Patent No. 6207442
                                                                                   Best Local
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LENGTH: 58
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APPLICANT: Sheppard, Paul O.
APPLICANT: Humes, Jacqueline M.
TITLE OF INVENTION: ADIPOCYTE-SPECIFIC PROTEIN HOMOLOGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 1998-10-15
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EARLIER APPLICATION NUMBER: 60/056,983
EARLIER FILING DATE: 1997-08-26
NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: 97-49
CURRENT APPLICATION NUMBER: US/09/140,804
CURRENT FILING DATE: 1998-08-26
                                                                                                                                                                  ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                               OTHER INFORMATION: synthetic oligonucleotide
                                                                                                                                                                                                   TYPE: DNA
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14
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o. 6197930
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                             1 AGATTTCTAGGAATTCAAATC 21
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AGAATACTAGGAATTCTACTC 34
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                                                                 Conservative
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                                                                               69.5%;
81.0%;
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81.0%;
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                                                                               Score 14.6; DB 3;
Pred. No. 4.2e+02;
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Pred. No. 4.2e+02;
                                                                 Mismatches
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RESULT 7 US-09-209-525-42

; Sequence 42,

Application US/09209525

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APPLICANT: Lok, Si
APPLICANT: Conklin, Darrell C.
APPLICANT: Conklin, Darrell C.
APPLICANT: Parrish, Julia E.
TITLE OF INVENTION: Mammalian Alpha Helical Protein-1
FILE REFERENCE: 97-71
FILE REFERENCE: 97-71
CURRENT APPLICATION NUMBER: US/09/209,525
CURRENT FILING DATE: 1998-12-10
NUMBER OF SEQ ID NOS: 56
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 42
LENGTH: 58
FYNE THAN
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Best Local Similarity
Matches 17; Conserv
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GENERAL INFORMATION:
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SOFTWARE: FastSEQ for
SEQ ID NO 46
LENGTH: 58
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                                                                                                                                                                             Sequence 35, Application US/09662402A Patent No. 6420117
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Best Local 9
                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sheppard, Paul O.
APPLICANT: Humes, Jacqueline M.
TITLE OF INVENTION: Adipocyte-Specific Protein Homologs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 97-49D1
CURRENT APPLICATION NUMBER: US/09/686,838B
CURRENT FILING DATE: 2000-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/140,804 PRIOR FILING DATE: 1998-08-26
              APPLICANT: Wessler, Susan R
APPLICANT: Casa, Alexandra M
TITLE OF INVENTION: MINIATURE INVERTED REPEAT TRANSPOSABLE ELEMENTS AND
TITLE OF INVENTION: METHODS OF USE
TILE REFERENCE: 235.00230101
CURRENT APPLICATION NUMBER: US/09/662,402A
CURRENT FILING DATE: 2000-09-14
CURRENT FILING DATE: 2000-09-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1997-08-26
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PRIOR APPLICATION NUMBER: 60/153,812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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Pred. No. 4.
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Pred. No. 4.2e+02;
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; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: Oligonucleotide primer US-09-662-402A-35
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LENGTH: 20
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NUMBER OF SEQ ID NOS: 40
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                                                        Matches
                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL
                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/20
APPLICATION UNBER: US/08/20
FILING DATE: 25-FEB-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryb A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                      TELEFAX: (619) 535-89-
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Wiggins, APPLICANT: Thomas, 1
                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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STATE: California
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                                                                                                                                            STRANDEDNESS:
                                                                                                                                 TOPOLOGY:
                                                                                                                                                              TYPE:
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16; Conservative
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AGTCCTAGGAATTCAAA 1
                           ATTTCTAGGAATTCAAA 19
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                                                         Conservative
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84.2%;
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Mammalian Glomerular Epithelial Protein
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88.2%;
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Pred. No. 6e+02;
0; Mismatches
                                                           0;
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                                                                         Score 13.8; DB 1
Pred. No. 9.4e+02
                                                           Mismatches
                                                                                                                                                                                                                                                                   9783
                                                                                        DB 1;
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                                                                                        Length 30;
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RESULT 11 US-08-463-090B-24/c

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RESULT 12
US-08-332-420-54/c
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                                                                                                                                                                                                                                                                                                                                                 Sequence 54, Application US/08332420 Patent No. 5744300
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                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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TITLE OF INVENTION: Humm
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                              APPLICANT: Maarten H.K. Linskens, et al.
TITLE OF INVENTION: METHODS AND REAGENTS FOR THE
TITLE OF INVENTION: IDENTIFICATION AND REGULATION
TITLE OF INVENTION: OF SENESCENCE-RELATED GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (617) 832-1299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cottarel, Guillaume
APPLICANT: Damagnez, Veronique
APPLICANT: Draetta, Guilio
TITLE OF INVENTION: Cell-Cycle Regulatory Proteins from
                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1)
                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: MIV032.01
                                                                                                                    COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/463,090B
FILING DATE: 05-JUN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII (text)
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                                                                                                                                                                                             ADDRESSEE:
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                                                                                                                                                       Los Angeles
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                                                                                                                                                                         E: Lyon & Lyon
633 West Fifth Street
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One Post Office Square
                                                                                                                      USA
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linear
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88.2%;
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Pred. No. 9.4e+02;
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APPLICANT: Lee, St
APPLICANT: Luckow,
APPLICANT: Leusch,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1,
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                                                                                             TELEFAX: (314)694-900 INFORMATION FOR SEQ ID NO:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                             SEQUENCE CHARACTERISTICS:
LENGTH: 37 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                         OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                          TELEPHONE: (314)694-5402
                                                                                                                                                                                                          APPLICATION NUMBER: US/0:
FILING DATE: 19920904
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Leusch, Mark S.
APPLICANT: Olins, Peter O.
TITLE OF INVENTION: Method
TITLE OF INVENTION: Eukary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/235,180
FILING DATE: April 29, 1994
ATTORNEY/AGENT INFORMATION:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                           STRANDEDNESS:
                                                                                                                                                                             NAME: Bennett, Dennis REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                               ZIP:
                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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STRANDEDNESS: single
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REGISTRATION NUMBER: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/332,420
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: Missouri
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Similarity 80.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Dennis A. Bennett, Monsanto Co., A3SG 800 N. Lindbergh Blvd.
                                                                                                            (314)694-9009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lee, Stephen C.
Luckow, Verne A.
                                                                                                                                                                                                                                                                                                             IBM PC compatible SYSTEM: PC-DOS/MS-DOS
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(genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Method of Producing Recombinant
Eukaryotic Viruses in Bacteria
                                                                                                                                                                                                                                                           US/07/941,363
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BER: 210/041
                                                                                                                                                                          34,547
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                                                                                                                                                              07-21 (872) A
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Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                            Version
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RESULT 14
US-09-143-634-28/c
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APPLICANT: Zdanovsky, Alexey G.
APPLICANT: Zdanovsky, Alexey G.
TITLE OF INVENTION: EXPRESSION OF CLOSTRIDIAL TOXINS AND PROTEINS
FILE REFERENCE: REMG-03193
CURRENT FILING DATE: 1998-08-28
CURRENT FILING DATE: 1998-08-28
NUMBER OF SEQ ID NOS: 28
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Best Local Similarity

Matches 16; Conserv
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SEQ ID NO 28
LENGTH: 38
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Patent No. 6214602
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OFERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                    APPLICATION NUMBER: US/08/308,196A FILLING DATE: 09-SEPT-1994 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07/983,523
                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Brown, Ma
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                   IIILE OF INVENTION: Production of Insulin-Like Growth
IIILE OF INVENTION: Factor-1 in Methylotrophic Yeast Cells
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
APPLICATION NUMBER:
                                   FILING DATE:
                                                                                                                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                ZIP:
                                                                                                                                                                                                                                                                                   CITY:
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: California
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                                                                                                                                                                                                                                                                                                   1660 Union Street
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                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                    Howard, Bradley D.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Davis, Geneva R.
Holtz, Gregory C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      Gleeson, Martin A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brierley, Russell A.
                                     03-MAR-1993
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80.0%;
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                                                                                                                                                                                                                                                                                                                 Martin,
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     US 07/578,728
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Pred. No. 1.
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Pred. No. 1.2e+03;
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                                                                                                                                             Version
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                                                                                                                                                                                                 PCT-US91-06452-4
                                                                                                                                                                                                                 RESULT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fort, Thomas L.
APPLICANT: McMillian, Ray A.
APPLICANT: Hellyer, Tobin
APPLICANT: You, Qimin
TITLE OF INVENTION: Amplification and Detection of Shiga-Like Toxin
TITLE OF INVENTION: Producing Organsims
FILE REFERENCE: SUT-1 Application
CURRENT APPLICATION NUMBER: US/09/289,803
CURRENT FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.0
SEC ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Fort, Th
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                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:Primer for SDA OTHER INFORMATION: of SLT-I Producing Organisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 51875
TELECOMMUNICATION INFORMATION:
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                             APPLICANT: Davis, Geneva R.
APPLICANT: Holtz, Gregory C.
APPLICANT: Gleeson, Martin A.
APPLICANT: Bradley, D. H.
TITLE OF INVENTION: Production of Insulin-Like Growth
TITLE OF INVENTION: Factor-1 in Methylotrophic Yeast Cells
                                                                                                                                           APPLICANT:
CORRESPONDENCE ADDRESS:
                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                              Application PC/TUS9106452
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                                                                                                                                           Brierley, Russell A.
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80.0%;
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Pred. No. 1.2e+03;
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US-09-289-803-2
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                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Applic Patent No. 6162605
                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Fort, Thomas L. APPLICANT: McMillian, Ray APPLICANT: Hellyer, Tobin
                                                                                                                                                                                                                                                APPLICANT: Hellyer, Tobin
APPLICANT: You, Qimin
TITLE OF INVENTION: Amplification and Detection of Shiga-Like Toxin
TITLE OF INVENTION: Producing Organsims
FILE REFERENCE: SLT-I Application
CURRENT APPLICATION NUMBER: US/09/289,803
CURRENT FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                            OTHER INFORMATION: Description of Artificial Sequence:Primer for OTHER INFORMATION: of SLT-I Producing Organisms
                                                                                                                                                      FEATURE:
                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                  LENGTH: 41
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LENGTH: 40 base pairs
TYPE: NUCLEIC ACID
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/578,728
FILING DATE: 04-SEP-1990
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Seidman, Stephanie L. REGISTRATION NUMBER: 33,779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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AGATTTCTAGGAATTCAAAT 20
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135 South LaSalle Street, Suite 900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (619)552-0095
                                 Conservative
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                                               Score 13.6; DB 3; Length 41; Pred. No. 1.2e+03;
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Pred. No. 1.2e+03;
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RESULT 20
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US-09-030-613-30
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US-09-289-803-1
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                                                                                                                 ; Sequence 30, Application US/09030613; Patent No. 6083706; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Patent No. 5221737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BARTSCH, KLAUS; SCHULZ, ARNO; UHLMANN, EUGEN TITLE OF INVENTION: GENE AND GENE STRUCTURE CODING FOR AMINOTRANSFERASE, AND MICROORGANISMS WHICH EXPRESS THIS G
                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:7:
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Patent No. 6102000.

GENERAL INFORMATION:
APPLICANT: Fort, Thomas L.
APPLICANT: McMillian, Ray A.
                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
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Patent No. 6162605
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APPLICANT: Hellyer, Tobin
APPLICANT: Hellyer, Tobin
APPLICANT: You, Qimin
TITLE OF INVENTION: Amplification and Detection of Shiga-Like Toxin
TITLE OF INVENTION: Producing Organsims
FILE REFERENCE: SIT-I Application
CURRENT APPLICATION NUMBER: US/09/289,803
CURRENT APPLICATION NUMBER: US/09/289,803
CURRENT FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 10
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
              CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
                                                                  APPLICANT: Florkiewicz, Robert Z. APPLICANT: Baird, J. Andrew TITLE OF INVENTION: INHIBITORS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 12
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                  NUMBER OF SEQUENCES:
 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 68
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6300 Columbia Center,
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                                                                    INHIBITORS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64.8%;
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80.0%;
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Pred. No. 1.2e+03;
0; Mismatches 4;
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Pred. No. 1.2e+03;
 701 Fifth Avenue
                                                                      LEADERLESS PROTEIN EXPORT
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STATE: 1

98104

CITY:

Seattle

Washington

USA

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US-09-313-221A-77; Sequence 77, Application US/09313221A; Patent No. 6468743
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                                                     RESULT 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-030-613-30
                                                                                                                                                                                                                                           ; OTHER INFORMATION: PCR Primer US-09-451-905-30
                                                                                                                                                                        Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 30, Application US/09451905 Patent No. 6306613
                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/451,905
CURRENT FILING DATE: 1999-12-01
NUMBER OF SEQ ID NOS: 48
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Robert Z. Florkiewicz APPLICANT: Andrew Baird APPLICANT: Dale E. Warnock TITLE OF INVENTION: MODULATORS OF TITLE OF INVENTION: AND METHODS FILE REFERENCE: 200124.402C4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (206) 682-603
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                               LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 76
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: No. 6083706tenburg Ph.D., Carol REGISTRATION NUMBER: 39,317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
                                                                                                                                                                            14;
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                                                                                                                                                                                          Similarity
                                                                                                        CTAGGAATTCAGATC 15
                                                                                                                                      CTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                   FastSEQ for Windows Version 4.0
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                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      MODULATORS OF LEADERLESS PROTEIN EXPORT AND METHODS FOR IDENTIFYING AND USING THE SAME
                                                                                                                                                                                          63.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.8%;
93.3%;
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Pred. No. 1.4e+03;
0; Mismatches 1
                                                                                                                                                                                            Pred.
                                                                                                                                                                                                            Score 13.4; DB 4; Length 30;
                                                                                                                                                                            Mismatches
                                                                                                                                                                                              No.
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US-08-840-316-5
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CURRENT FILING DATE: 1999-05-17
PRIOR APPLICATION NUMBER: US 60/086,025
PRIOR FILING DATE: 1998-05-18
NUMBER OF SEQ ID NOS: 145
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/08840316
Patent No. 6054567
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APPLICANT: Mark S. Fraser (Inventor)
TITLE OF INVENTION: PCR TECHNIQUES FOR DETECTING MICROBIAL
TITLE OF INVENTION: AND VIRAL CONTAMINANTS IN FOODSTUFFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Richard W. Bork
REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 2026
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                          TELEFAX: (212) 751-684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 40
                                                                                                                                                                                                                                                                                                                            FILING DATE: 11-APR-CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOCTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Emerson, APPLICANT: Tsarev,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 345 PARK AVENUE CITY: NEW YORK
                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE:
                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                      LENGTH:
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                                                                        25 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Emerson, Suzanne U., Purcell, Robert H.,
Tgarev, Sergel. A., and Robinson, Robin A.
VENTION: Recombinant Proteins Of
VENTION: A Pakistani Strain Of Hepatitis E And Their
VENTION: Use In Diagnostic Methods And Vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HUNT-042784
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                                                                                                                                                                                                                                                                                                                                                                UMBER: US/08/840,316
11-APR-1997
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Pred. No. 1.4e+03;
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0;

Query Match Best Local

Match 62.9%; Score 13.2; DB 3; Local Similarity 83.3%; Pred. No. 1.7e+03;

Length 25;

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RESULT 26
US-08-471-971-5/c
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            Sequence 5, Application US/08471971 Patent No. 6287759
                                                                                                                                                                   Query Match
Best Local (
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                                                                                                                                                          Matches
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Tsarev, Sergei. A., 1
APPLICANT: Suzanne U., Purcell,
                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 18-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Richard W. Bork
REGISTRATION NUMBER: 36,4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 20.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: USO8, FILING DATE: 03-OCT-1994 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 03-OCT-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-MAY-1997 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                            TELEPHONE: (212)
                                                                                                                                                                                                                          STRANDEDNESS: SIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                     LENGTH:
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                                                                                                                                                                      Similarity
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                                                                                                                           ATTICTAGGAATTCAAAT 20
                                                                                                                                                                                                                                                        nucleic acid
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                                                                                                                                                                                                                                                                     25 base pairs
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                                                                                                                                                                                                                                          single
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7 751-6849
7 NO: 5:
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" Purcell, Robert H.
                                                                                                                                                                     62.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Recombinant Proteins Of
A Pakistani Strain Of Hepatitis E And Their
Use In Diagnostic Methods And Vaccines
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                                                                                                                                                      ; Score 13.2; D; Pred. No. 1.7e
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                                                                                                                                                                     .7e+03;
                                                                                                                                                                                 DB 3;
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                                                                                                                                                                                     Sequence 5, Application US/09402776
Patent No. 6458562
GENERAL INFORMATION:
APPLICANT: Emerson, Suzanne U.,
APPLICANT: Tsarev, Sergei. A., 6
                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                     APPLICANT: Emerson, Suzanne U., Purcell, Robert H., APPLICANT: Tearev, Sergei. A., and Robinson, Robin A.
TITLE OF INVENTION: Recombinant Proteins Of
TITLE OF INVENTION: A Pakistani Strain Of Hepatitis E And Their
TITLE OF INVENTION: Use In Diagnostic Methods And Vaccinc-
NUMBER OF SEQUENCES: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435

PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US08/316,765

FILING DATE: 03-OCT-1994

CLASSIFICATION: 435

PRIOR APPLICATION DATA:
                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Richard W. Bork
REGISTRATION NUMBER: 36,459
REFERENCE DOCKET NUMBER: 20)
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                           ADDRESSEE: MC
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COUNTRY: U
ZIP: 10154
                             CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US07/947,263
FILING DATE: 18-SEP-1992
CLASSIFICATION: 435
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TELEFAX: (212) 75
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                                                                                                                                                                                                                                                                                                                                                                                                                       ch 62.9%;
l Similarity 83.3%;
15; Conservative
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                                                                PARK AVENUE
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/ 751-6849
- MO: 5:
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A Pakistani Strain Of Hepatitis E And Their
Use In Diagnostic Methods And Vaccines
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Pred. No. 1.7e+03;
0; Mismatches 3;
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COMPUTER READABLE FORM

MEDIUM TYPE:

IBM PC COMPATIBLE FLOPPY DISK

PC-DOS/MS-DOS

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RESULT 28
PCT-US93-08849A-5/c
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APPLICANT:
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TELEFAX: (212) 751-6849
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                   SOFTWARE: WORDPERFECT 5.1

CURRENT APPLICATION DATA:

APPLICATION UMBER: PCT/US93/08849.

FILING DATE: 17-SEP-1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US07/947,263

FILING DATE: 18-SEP-1992

ATTORNEY/AGENT INFORMATION:

ANDER: 18-SEP-1992

ATTORNEY/AGENT INFORMATION:
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                                  TELEFAX: (212) 751-6849
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION NUMBER:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
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TITLE OF INVENTION:
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REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
                                                  REFERENCE/DOCKET NUMBER: 2026-4032 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
ZIP: 101
                                                                                                                       NAME: William S. Feiler REGISTRATION NUMBER: 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
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STREET: 34
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83.3%;
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A Pakistani Strain Of Hepatitis E And Their
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Pred. No. 1.7e
0; Mismatches
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RESULT 30 US-09-182-816-30/c

Sequence 30, Application US/09182816 Patent No. 6143542 GENERAL INFORMATION:

APPLICANT: Wisnewski, Nancy

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PCT-US93-08849-5/c
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                                                          Matches
                                                                      Query Match
Best Local :
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                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/947,263
                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIB:
OPERATING SYSTEM: PC-DOS/I
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: FLOPPY
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ADDRESSEE: MORGAN & FINNEGAN
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                                                                                                                                             STRANDEDNESS: single
                                                                                                                                                                                                                                                                             NAME: Bork, Richard, W. REGISTRATION NUMBER: 36,459
                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/
FILING DATE: 17-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: NEW YORK
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ATGTCTGTGAATTCAAAT 3
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SYSTEM: PC-DOS/MS-DOS
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83.3%;
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83.3%;
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A Pakistani Strain Of Hepatitis E And Their
Use In Diagnostic Methods And Vaccines
98
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Pred. No. 1.
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                                                                        Score 13.2; DB 5;
Pred. No. 1.7e+03;
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                                                                                    Length 25;
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                                                           Indels
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RESULT 32
US-09-634-530-30/c
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US-09-471-528-30/c
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               Sequence 30, Application US/09634530 Patent No. 6290958 GENERAL INFORMATION:
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LENGTH: 31
                                                                                                                                                                                                                              Matches
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Best Local
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LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/471,528
CURRENT FILLING DATE: 1999-12-27
EARLIER APPLICATION NUMBER: 09/182,816
EARLIER FILLING DATE: 1998-10-29
EARLIER APPLICATION NUMBER: 08/989,510
EARLIER APPLICATION NUMBER: 08/989,510
EARLIER FILLING DATE: 1997-12-12
NUMBER OF SEQ ID NOS: 35
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lo, Katherine C.
APPLICANT: Brandt, Kevin S.
TITLE OF INVENTION: FILEA EPOXIDE HYDROLASE PROTEINS AND USES
FILE REFERENCE: FC-3-C1-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOVEL FLEA EPOXIDE HYDROLASE NUCLEIC ACID MOLECULES, TITLE OF INVENTION: PROTEINS AND USES THEREOF FILE REPERIORS: FC-3-C1 CURRENT APPLICATION NUMBER: US/09/182,816 CURRENT FILING DATE: 1998-10-29 EARLIER APPLICATION NUMBER: 08/989,510 EARLIER FILING DATE: 1997-12-12 NUMBER: 08/989,510 EARLIER FILING DATE: 1997-12-12 NUMBER: 08/989,510 EARLIER FILING DATE: 1997-12-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Silver, Gary M. APPLICANT: Lo, Katherine C. APPLICANT: Brandt, Kevin S.
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                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Descri
                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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Wisnewski,
                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                      Description of Artificial Sequence:
                                                                                                                                                                                                                                            62.9%;
83.3%;
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Pred. No. 1.
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Pred. No. 1.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: ; OTHER INFORMATION: US-09-634-530-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: FILA EPOX.
FILE REFERENCE: FC-3-C1-1
CUERENT APPLICATION ***-
CUERENT APPLICATION ***-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 6, Applic Patent No. 5360733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Tocal Similarity
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SEQ ID NO 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-12-27
PRIOR APPLICATION NUMBER: 099/
PRIOR FILING DATE: 1998-10-29
PRIOR APPLICATION NUMBER: 08/9
PRIOR FILING DATE: 1997-12-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/634,530
CURRENT FILING DATE: 2000-08-08
PRIOR APPLICATION NUMBER: 09/471,528
PRIOR FILING DATE: 1999-12-27
                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                  STREET: 43, STREET: SAN DIEGO CITY: SAN DIEGO CTATE: CALIFORNIA
                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN
REGISTRATION NUMBER: 31,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BIERHUIZEN, MARTI FA
TITLE OF INVENTION: A NOVEL BETAL-6
TITLE OF INVENTION: N-ACETYLGUCOSAMINYLTRANSFERASE, ITS
TITLE OF INVENTION: LEUKOSIALIN AND A METHOD FOR CLONING
TITLE OF INVENTION: ENZYMATIC ACTIVITY
                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
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                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                 FILING DATE: 1: CLASSIFICATION:
                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                TELEPHONE:
STRANDEDNESS:
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                 NUCLEIC
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                             33 base pairs
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                                                                                619-535-8949
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83.3%;
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Pred. No. 1.8e
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PROTEINS HAVING
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                                                                              US-08-472-482-6/c
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                                                                                                   RESULT 35
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Sequence 6, Application US/08472482
Patent No. 5658778
GENERAL INFORMATION:
APPLICANT: BIERHUIZEN, MARTI FA
APPLICANT: BIERHUIZEN, MARTI FA
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Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
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CORRESPONDENCE ADDRESS:
ADDRESSEE: CAMPBELL AND FLORES
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APPLICANT:
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LENGTH: 33 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 619-535-8949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/227,455
FILING DATE: 14-APR-1994
CLASSIFICATION: 435
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TOPOLOGY: linear
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NUENTION: A NOVEL BETA1-6
VENTION: A NOVEL BETA1-6
VENTION: N-ACETYLGLUCOSAMINYLTRANSFERASE, ITS ACCEPTOR MOLECULE,
VVENTION: LEUKOSIALIN AND A METHOD FOR CLONING PROTEINS HAVING
NVENTION: EXYMATIC ACTIVITY
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Pred. No. 1
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                                                                                                                                                                                                                                     Pred. No.
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US-08-487-069-6/c
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                                                                                                                                                                                                                                                                                                          Patent No. 5684134
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/955
FILING DATE: 01-OCT-1992
ATTORNEY, AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ
TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: US
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,482
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             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                           COUNTRY: U
ZIP: 92122
                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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SOFTWARE:
                                                                                                                              CITY: SAN DIEGO
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                                                                                                           CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: CAMPBELL AND FLORES
4370 LA JOLLA VILLAGE DRIVE, SUITE 700
                                                                                                                                           E: CAMPBELL AND FLORES
4370 LA JOLLA VILLAGE DRIVE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                        BIERHUIZEN, MARTI FA
VENTION: A NOVEL BETA1-6
VENTION: N-ACETYLGUUCOSAMINYLTRANSFERASE,
VENTION: LEUKOSIALIN AND A METHOD FOR CLOI
VENTION: ENZYMATIC ACTIVITY
                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                           FUKUDA,
PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     619-535-9001
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N-ACETYLGLUCOSAMINYLTRANSFERASE, ITS
LEUCOSIALIN AND A METHOD FOR CLONING
ENZYMATIC ACTIVITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62.9%;
                                                                                                                                                                                                                                                                          MARTI FA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07/955,041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P-LJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.2; DB 1;
Pred. No. 1.8e+03;
Version #1.25
                                                                                                                                               SUITE 700
                                                                                                                                                                                                                            RASE, ITS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 33;
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                                                                                                                                                                                                                              ACCEPTOR
PROTEINS
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RESULT 37
US-08-169-715-47/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; MOLECULE TYPE: cDNA US-08-487-069-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 47, Application US/08169715 Patent No. 6300056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 62.9%;
Best Local Similarity 83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                  TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 619-535-8949 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                            FILING DATE: 18-DEC-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: HIV PROBES FOR USE IN SOLUTION PHASE TITLE OF INVENTION: SANDWICH HYBRIDIZATION ASSAYS NUMBER OF SEQUENCES: 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Irvine, Bruce
APPLICANT: Horn, Thomas
APPLICANT: Chang, Chu-An
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 01-OCT-1992
ATTORNEY/AGENT INFORMATION:
               SEQUENCE CHARACTERISTICS:
                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 415-813-5600
                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                         NAME: Monroy, Gladys H.
REGISTRATION NUMBER: 32,430
REFERENCE/DOCKET NUMBER: 22300-20150.00
                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 545 Middlefield Road, Suite 200 CITY: Menlo Park
                                                                                                                                                                                                                                                       CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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REFERENCE/DOCKET NUMBER: P-LJ
                                                                     TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/955,041
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/487,069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               California
33 base pairs
                                                                       415-327-2951
                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bruce D.
                                                                                                                                                                                                                       07/813,583
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Pred. No. 1.8e+03;
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                                                  ; CTHER INFORMATION: Oligonucleotide ZC15223 US-09-686-838B-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Oligonucleotide ZC15223 US-09-140-804-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-140-804-38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-686-838B-38
                                                                                                                                                        NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 38
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 38, Application US/09686838B Patent No. 6482612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 47
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 38
LENGTH: 63
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Query Match
Best Local Similarity
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APPLICANT: Humes, Jacqueline M.
TITLE OF INVENTION: ADIPOCYTE-SPECIFIC PROTEIN HOMOLOGS
FILE REFERENCE: 97-49
CURRENT APPLICATION NUMBER: US/09/140,804
CURRENT FILING DATE: 1998-08-26
EARLIER APPLICATION NUMBER: 60/056,983
EARLIER FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/686,838B CURRENT FILING DATE: 2000-10-10 PRIOR APPLICATION NUMBER: US 09/140,804
                                                                                                                                                                                                                                                                                                                         APPLICANT: Sheppard, Paul O.
APPLICANT: Humes, Jacqueline M.
TITLE OF INVENTION: Adipocyte-Specific Protein Homologs
FILE REFERENCE: 97-49D1
                                                                                                                                                                                                              PRIOR FILING DATE: 1998-08-26
PRIOR APPLICATION NUMBER: US 60/056,983
PRIOR FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                    ORGANISM: Artificial Sequence FEATURE:
                                                                                                                          TYPE: DNA
                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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Similarity 83.3%;
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62.9%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 13.2; DB 3;
Pred. No. 1.8e+03;
0; Mismatches 3
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Pred. No. 1.8e+03;
Score 13.2; DB 4;
Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 3;
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               Length 63;
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COUNTEX:

ZIP: 10036

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/413,118

FILING DATE: 29-MAR-1995

CLASSIFICATION NUMBER: US 08/220,151

FRIING DATE: 30-MAR-1994

ATTORNEY/AGENT INFORMATION:

NAME: FROWMER, WILLIAM S.

REGISTRATION NUMBER: 25,506

REFERENCE/MOUNICATION INFORMATION:

TELEPHONE: (212) 840-333

TELEPHONE: (212) 840-333

TELEPHONE: (212) 840-333

TELEPHONE: (212) 840-312:

SEQUENCE CHARROTERISTICS:

SEQUENCE CHARROTERISTICS:

LENGTH: 72 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: CDNA

US-08-413-118-122
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Search completed: February 4, 2004, 17:53:49 Job time: 52.4091 secs
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US-08-413-118-122/c
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Patent No. 5688920

GENERAL INFORMATION:
APPLICANT: PAOLETTI, ENZO
APPLICANT: LIMBACH, KEITH J.
TITLE OF INVENTION: NUCLEOTIDE AND AMINO ACID SEQUENCES OF
TITLE OF INVENTION: CANINE HERPESVIRUS GB, GC, AND GD AND USES THEREFOR
NUMBER OF SEQUENCES: 128
                                                                                                                                                                    Query Match 62.9
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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ADDRESSEE: CURTIS, MORRIS & SAFFORD, P.C.
STREET: 530 FIFTH AVENUE, 25TH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: NEW YORK STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                46 AGAATACTAGGAATTCTA 63
                                                                                     18 AAATGTTTAGGAATTCAA 1
                                                                                                             1 AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AGATTTCTAGGAATTCAA 18
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                                                                                                                                                               62.9%; Score 13.2; DB 1; Length 72; 83.3%; Pred. No. 1.8e+03; ative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Therapeutic agent for treating transcription factor-related illnesses such as proliferative malignancies, comprises an oligonucleotide for regulating transcription factor function -
                                                                                         Antigenic composition comprises polynucleotide fragment - used as vaccine to treat or prevent e.g. cancer or pat infections and to modulate immune response e.g. toleranc regulation of THI/TH2 cells
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neoplastic diseases, and immunological and inflammatory disorders.
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TLR7 and TLR8 polypeptides. These polypeptides comprise fully defined sequences of 1032, 1050 or 1032 amino acids as given in specification, their fragments, where TLR9, TLR7 and TLR8 polypeptides or their fragments have an amino acid sequence which is identical to human TLR9, TLR7 or TLR8 polypeptides or their fragment except for at least one amino acid of a murine TLR polypeptide. The isolated nucleic acids of the invention are useful for inhibiting TLR9 signalling activity in a cell. TLR7, TLR8 and TLR9 polypeptides are useful for identifying nucleic acid molecules which interact with a TLR polypeptide or its fragment. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             response and the modulation is selected from the group break of tolerance, regulation of TH1/TH2 helper cell responses, switch of Ig classes, treatment of autoimmune responses and induction of tolerances. DNA oligomers are used to enhance the reactivity of immune cells to viral, bacterial and parasitic antigens, to break tolerance in anergic and B cells e.g. against tumour antigens, as adjuvants in vaccination against tumour-defined antigens and immunostimulatory substances in an immune response against tumours and to suppress immune reactions of the innate and acquired immune system. The composition is inexpensive and stable and does not cause lethal shock, which happens with prior art
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated murine Toll-like receptor (TLR)9, TLR7, TLR8 polypeptides, useful for identifying species specificity of immunostimulatory nucleic acid and identifying immunostimulatory nucleic acids -
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                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 76; 195pp; English
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23-JAN-2001; 2001US-263659
17-MAY-2001; 2001US-291726P
22-JUN-2001; 2001US-300210P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (COLE-)
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                                                                                                                                                                                                                                                                                                                        invention relates to isolated murine
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08-JUN-2001;
08-JUN-2001;
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                         transcription factor-protein interactions. The methods of the invention are also useful for facilitating the screening and identification of new drugs, characterising their mechanism of action and screening for adverse side effects based on drug's impact expression. The present DNA sequence represents a probe used in the method of the invention.
                                                                                                    The invention comprises a method for identifying complexes between a transcription factor and another protein. The invention also comprises method for isolating DNA probes which bind to activated transcription factors. The methods of the invention are useful for identifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying transcription factor-protein complexes, by isolating transcription factor complexes from sample based on a specific type factor, and identifying different proteins present in isolated
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2001US-0877738.
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                                                                                                                                                                                                                                                                                                                                 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                    other;
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72;
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72;
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ABT17327 standard; DNA; 63 BP

spieen; human;

primer; ss

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RESULT 7
AAA547
XX
AC AAA5
XC AAA5
XX
DT 11-A
XX
DE DNA
XX
KW zcyt
KW zcyt
KW baind
KW haem
KW immu
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                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                            The invention comprises a method for identifying complexes between a transcription factor and another protesin. The invention also comprises a method for isolating DNA probes which bind to activated transcription factors. The methods of the invention are useful for identifying transcription factor-protein interactions. The methods of the invention are also useful for facilitating the screening and identification of new drugs, characterising their mechanism of action and screening for adverse side effects based on drug's impact expression. The present DNA sequence represents a probe used in the method of the invention.
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08-JUN-2001; 2001US-0877705.

08-JUN-2001; 2001US-0877738.

05-SEP-2001; 2001US-0947274.
             zcytor 10 cytokine receptor; cytokine; receptor; antibody; ligand; binding; detection; modulation; recombinant cell; haematopoietic cell; lymphoid cell; myeloid cell; lymph; immune system; blood; bone; inflammatory response; inflammation;
                                                                                                               11-APR-2001
                                                                                                                                                                     AAA54479 standard;
                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-148829/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-JUN-2001; 2001US-0877243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2002; 2002WO-US17408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Probe; ss; transcription factor-protein complex; transcription factor; drug screening; drug identification; array hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Identifying transcription factor-protein complexes, by isolating transcription factor complexes from sample based on a specific tyfactor, and identifying different proteins present in isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PANO-)
                                                                                                                                                                                                                                        63
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                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                  63 BP; 21 A; 9 C; 12 G; 21 T; 0 other;
                                                                                                                                                                                                                                                                   AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                                                         AGATTTCTAGGAATTCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fig
                                                                                                                                                                                                                                                                                             Conservative
                                                                                    comprising
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                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6; 167pp;
                                                                                                                                                                      DNA; 100
                                                                                                                                                                                                                                                                                                                          85.7%;
                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                     STAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
                                                                                                                                                                                                                                         46
                                                                                                                                                                      ВÞ
                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                            Score 18;
Pred. No.
                                                                                    transcription factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         array hybridisation
                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                            DB
75
                                                                                                                                                                                                                                                                                                                          25;
                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                       Length 63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              type
                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                    AAA54480/c
                                                                                                                                                                                                                                  RESULT 8
                                                                                                                                                                                                                                                                                                                                                                                                                                              of zcytor10 receptor ligand within a test sample can be achieved.
The method comprising contacting a test sample containing an amino acid sequence from Cys15 or Gly25 to Pro230 of the zcytor 10 cytokine receptor and detecting the binding of the polypeptide to a C ligand in the sample. Specified peptide fragments of the zcytor 10 cytokine receptor and the methods described are used to identify C ligands that stimulate the proliferation and/or development of the cytokine receptor are useful for treating lymphoid, immune, c the cytokine receptor are useful for treating lymphoid, immune, c inflammatory, splenic, blood or bone disorders and for generating antibodies directed against the receptor. An exemplary luciferase mammalian expression vector is the KZ134 plasmid which was constructed with two complementary oligonucleotides (AAA5449), which comprise STAT transcription factors from 4 genes (a modified c-fos Sis element, the p21 SIE1 from the p21 WAPI gene, the mammary gland response element of the Beta-casein gene
                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                      Best Local
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
            zcytor 10 cytokine receptor; cytokine; receptor; antibod binding; detection; medulation; recombinant cell; haematopoietic cell; lymphoid cell; myeloid cell; lymph; immune system; blood; bone; inflammatory response; infla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor. Those cells can then be used to detect the presence of a modulator of zcytor10 protein by culturing the cells in the presence of a test ligand and comparing levels of activity of mouse zcytor10 in the presence of the test manual absence of the test of activity of detection of zcytor10 receptor ligand within a test manual. Similarly, detection
 spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cytokine receptor mouse zcytor 10, useful for detecting ligands that stimulate proliferation or development of haematopoietic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAY-2000; 2000WO-US12924
                                                                                                   DNA fragment
                                                                                                                                  11-APR-2001
                                                                                                                                                                                                   AAA54480
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolating a nucleotide which encodes the zcytor 10 cytokine receptor enables the production of recombinant cells expres
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 19; Page 128; 134pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-016096/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                     67
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                                                                                                                                                                                                                                                                                                                                                     18;
human; primer; ss
                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       100
                                                                                                                                                                                                                                                                                                      AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                     AGATTTCTAGGAATTCAA
                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 24 A; 33 C; 17 G; 26 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              myeloid
                                                                                               comprising STAT transcription factors
                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0309861
                                                                                                                                                                                                 DNA; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cells
                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                    85.7%;
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                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                     84
                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                      Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English
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                                                                receptor; antibody; ligand;
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                                                                                                                                                                                                                                                                                                                                                                                    22; Length 100;
                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                     Indels
                 inflammation,
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ABA93801
ID ABA
XX
AC ABA
AC ABA
AX
DT 01-XX
KW Zcy
KW Zcy
KW ant
KW ant
KW inff
KW aut

ABA93801; 01-MAY-2002 KZ134 plasmid

(first entry)

antiinflammatory; antiviral; antirheumatic; antiarthritic; cytostatic; muscular; lymphoid; immune; inflammatory; spleenic; blood; bone; infection; immunosuppression; cytotoxicity; leukopenia; Crohn's disease;

Zcytor17; chromosome 5; 5q11; cytokine receptor;

immunomodulatory;

NO:43.

construction oligonucleotide SEQ ID

autoimmune

disease;

rheumatoid arthritis;

muitipie

sclerosis; cancer;

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38

AGATTTCTAGGAATTCAA

21

0;

RESULT 9

ABA93801 standard;

UNA;

100

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                                                                                                                    CC receptor enables the production of recombinant cells expressing the creceptor. Those cells can then be used to detect the presence of a cc modulator of zcytor10 protein by culturing the cells in the presence of a cc modulator of zcytor10 protein by culturing the cells in the presence cc in the presence and absence of the test sample. Similarly, detection of zcytor10 receptor ligand within a test sample can be achieved. The method comprising contacting a test sample containing an amino cc acid sequence from Cysl5 or Gly25 to pro230 of the zcytor 10 cytokine receptor and detecting the binding of the zcytor 10 cytokine receptor and the methods described are used to identify cligands that stimulate the proliferation and/or development of haematopoietic, lymphoid and myeloid cells. Peptide fragments of the cytokine receptor are useful for treating lymphoid, immune, clift the cytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor are useful for treating lymphoid, immune, clift eytokine receptor sector is the KZ134 plasmid which was constructed with two complementary oligonucleotides (AAA54479, clift eytokine constructed with two complements of the Erg Pl reme beta-casein gene constructed constructed selement of the Erg Pl reme
                                      Best Loc
Matches
                                                                             Query Match
                                                                                                                    Sequence 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cytokine receptor mouse zcytor 10, useful for detecting ligands that stimulate proliferation or development of haematopoietic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-016096/02
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                                                             Local
                                                                                                                                                          ζij,
                                                                                                                                                        STAT inducible element of the Fcg
  Ц
                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19;
                                                             Similarity
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AGATTTCTAGGAATTCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 128;
                                        Conservative
                                                                                                                  BP; 26 A; 17 C; 33 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        myeloid cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-US12924
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                                                         100.0%;
                                                                             85.7%;
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  18
                                      0,
                                                           Score 18;
Pred. No.
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                                        Mismatches
                                                                                                                    24 T; 0
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                                                     DB
76;
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                                                                             22;
                                                                                                                                                        gene.
                                                                                                                    other;
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                                        Indels
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ABA93802/c
ID ABA938
XX
AC ABA938
XX
DT 01-MAY
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DT 02-MAY
XX
Cytor
KW Zcytor
KW Zcytor
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KW infect
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                                                                                                                                                                                                                                                                    RESULT 10
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunity and for stimulating lymphocyte proliferation, such as in the treatment of infections involving immunosuppression, including certain viral infections. They are also useful for inducting cytotoxicity and for treating leukopenias. Antagonist of zcytor17 polypeptides are useful for treating autoimmune diseases (e.g. rheumatoid arthritis and multiple sclerosis), inflammatory diseases (e.g. Crohn's disease), cancer, pancreatitis, and inflammatory bowel disease. Zcytor17 was mapped to chromosome 5, specifically to the 5q11 chromosomal region. ABA93767 to ABA93843 and ABB05730 to ABB05745 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zcytor17 has immunomodulatory, antiinflammatory, antiviral, cytostatic, antirheumatic, antiarthritic and muscular activities. The zcytor17 proteins are useful for treating and diagnosing lymphoid, immune, inflammatory, spleenic, blood or bone disorders. Agonists or anti-zcytor17 antibodies are useful in stimulating cell-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful for treating and diagnosing lymphoid, spleenic, blood or bone disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sprecher CA,
Maurer MF;
                                infection; immunosuppression; cytotoxicity; leukopenia; Crohn's disease;
autoimmune disease; rheumatoid arthritis; multiple sclerosis; cancer;
                                                               antiinflammatory; antiviral; antirheumatic; antiarthritic; cytostatic;
muscular; lymphoid; immune; inflammatory; spleenic; blood; bone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
PCR primer; probe; ss
                                                                                                    Zcytor17; chromosome
                                                                                                                                                                    01-MAY-2002
                                                                                                                                                                                                   ABA93802;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer; probe;
                                                                                                                                 KZ134 plasmid construction oligonucleotide SEQ ID NO:44.
                                                                                                                                                                                                                                    ABA93802 standard; DNA; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ZYMO ) ZYMOGENETICS
                   inflammatory disease; pancreatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nflammatory disease; pancreatitis; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                     67
                                                                                                                                                                                                                                                                                                                                                                                     18;
                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                     100
                                                                                                                                                                                                                                                                                                                                                  AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                                                                                                                                     AGATTTCTAGGAATTCAA 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention describes a cytokine receptor designated zcytor17.
                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 24 A; 33 C; 17 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001US-267963P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-214282P.
2000US-214955P.
                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Presnell
                                                                                                                                                                                                                                                                                                                                                                                                   85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INC
                                                                                                5; 5q11; cytokine receptor; immunomodulatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SR,
                                                                                                                                                                                                                                    ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gao
                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                   Score 18;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   26
                   inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Whitmore TE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   T;
                                                                                                                                                                                                                                                                                                                                                                                                 DB 24; Length 100; 76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immune,
                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kuijper JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       zcytor17 which inflammatory,
                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 11
AAS20691
ID AAS20
XX AAS20
XX AAS20
XX Plasn
DT 09-AE
XX Cytob
KW natus
KW B-cel
KW B-cel
KW inmus
XX immus
XX Syntl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a cytokine receptor designated zcytor17. CC Zcytor17 has immunomodulatory, antiinflammatory, antiviral, cytostatic, cc antirheumatic, antiarthritic and muscular activities. The zcytor17 cc antirheumatic, antiarthritic and muscular activities. The zcytor17 cc proteins are useful for treating and diagnosing lymphoid, immune, inflammatory, spleenic, blood or bone disorders. Agonists or canti-zcytor17 antibodies are useful in stimulating cell-mediated cc immunity and for stimulating lymphocyte proliferation, such as in the treatment of infections involving immunosuppression, including certain cc treatment of infections. They are also useful for inducting cytotoxicity and coveral infections. They are also useful for inducting cytotoxicity and creating autoimmune diseases (e.g. rheumatoid arthritis and multiple csclerosis), inflammatory diseases (e.g. Crohn's disease), cancer, cc pancreatitis, and inflammatory bowel disease. Zcytor17 was mapped to chromosome 5, specifically to the 5q11 chromosomal region. ABA93767 to ABA93843 and ABB05730 to ABB05745 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                             Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; T-cell proliferation; B-cell proliferation; anti-tumour response; immune system; immunostimulant; cytostatic; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated polynucleotide encoding a cytokine useful for treating and diagnosing lymphoid, spleenic, blood or bone disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JUN-2000; 2000US-214282P.
29-JUN-2000; 2000US-214955P.
08-FEB-2001; 2001US-267963P.
                                                                                                                                                                                               09-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 100 BP; 26 A; 17 C; 33 G; 24 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sprecher CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-JUN-2001; 2001WO-US20484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200200721-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                               Synthetic.
                                                                                                                                                             Plasmid KZ 134 oligonucleotide
                                                                                                                                                                                                                                                             AAS20691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ZYMO ) ZYMOGENETICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                           18;
                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                        AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                                                                                                                                                             AGATTTCTAGGAATTCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 190;
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Presnell SR,
                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                         85.7%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    235pp; English.
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                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                          °;
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 18;
Pred. No.
                                                                                                                                                                ZC12749.
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                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Whitmore
                                                                                                                                                                                                                                                                                                                                                                                                                           DB
76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Œ,
                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 100;
                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kuijper JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     zcytor17 which
inflammatory,
                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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US6307024-B1

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RESULT 12
AAS20692/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The zalphall Ligand polypeptide is also useful in preparing antibodies that bind to zalphall Ligand epicopes. The zalphall Ligand polynucleotides can be used as probes or primers to clone regions of a zalphall Ligand gene, and in gene therapy. Zalphall Ligand may also be used to identify inhibitors of its activity, to enhance the generation of anti-tumour responses with or without the infusion of donor lymphocytes, and to activate or stimulate the immune system. The present sequence represents an oligonucleotide used to construct plasmid XZ 134 in the methods of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         zalphall Ligand and the polynucleotide encoding it. The invention also gives the sequence for the zalphall receptor and the polynucleotide encoding it. The zalphall Ligand polypeptide stimulates proliferation of natural killer (NK) cells or NK cell progenitors, the activation of NK cells, proliferation of T-cells, proliferation of B-cells stimulated with anti-CD40 antibodies, stimulates an antigenic response in a mammal, and reduces proliferation of B-cells stimulated with anti-ID40 antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-MAR-1999;
11-MAR-1999;
01-JUL-1999;
                                                                                                                                    natural killer cell proliferation; B-cell proliferation; anti-tumour x
                                                                                                                                                 Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; T-cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New zalphall ligand polypeptides and polynucleotides, useful for stimulating proliferation, activation, differentiation and/or induction of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antigenic response
09-MAR-2000; 2000US-0522217
                              23-OCT-2001.
                                                                                                                      immunostimulant; cytostatic;
                                                                                                                                                                                                  Plasmid KZ
                                                                                                                                                                                                                                                                                            AAS20692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                       67
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                                                                                                                                                                                                                                                                                                                                                                                                                                  18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100
                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                               AGATTTCTAGGAATTCAA 18
                                                                                                                                                                                                                                                                                                                                                                       AGATTTCTAGGAATTCAA 84
                                                                                                                                                                                               134 oligonucleotide ZC12748.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Presnell SR,
V, Nelson AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Column 149-150; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000US-0522217
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99US-123904P.
99US-142013P.
                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 32 C; 17 G; 26 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              85.7%;
                                                                                                                                      anti-tumour response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      relates to the isolation of a novel cytokine, he polynucleotide encoding it. The invention
                                                                                                                                                                                                                                                                                            100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sprecher CA,
Dillon SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18;
Pred. No
                                                                                                                    primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                  No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Foster DC, Hammond AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 100;
                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cc zalphall Ligand and the polynucleotide encoding it. The invention calso gives the sequence for the zalphall receptor and the polynucleotide encoding it. The zalphall Ligand polypeptide stimulates proliferation of cells progenitors, the activation of natural killer (NK) cells or NK cell progenitors, the activation of NK cells, proliferation of E-cells, proliferation of B-cells stimulated with anti-CD40 antibodies, stimulates an antigenic response in a mammal, and reduces proliferation of B-cells stimulated with anti-Lyd antibodies. The zalphall Ligand polypeptide is also useful in preparing antibodies. The zalphall Ligand polypeptide is also useful in preparing antibodies chat bind to zalphall Ligand epitopes. The zalphall Ligand may construct polynucleotides can be used as probes or primers to clone regions of a zalphall Ligand gene, and in gene therapy. Zalphall Ligand may also be used to identify inhibitors of its activity, to enhance the generation of anti-tumour responses with or without the infusion of construct land to activate or stimulate the immune system.

The present sequence represents an oligonucleotide used to construct plasmid KZ 134 in the methods of the present invention.
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Matches
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11-MAR-1999;
01-JUL-1999;
                                                                                                                                                                                  inflammatory disorder; cell proliferation; immune disorder; cancer; SL systemic lupus erythematosus; myasthenia gravis; rheumatoid arthritis; diabetes; autoimmune disease; multiple sclerosis; ulcerative colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 100 BP; 26 A; 17 C; 32 G; 25 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New zalphall ligand polypeptides and polynucleotides, useful for
stimulating proliferation, activation, differentiation and/or induction
of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                      asthma;
                                                                                                                                                                                                                                      Zalphall; cytokine receptor; immunosuppressive; cytostatic; haemostatic;
                                                                                                                                                                                                                                                                           BaF3/KZ134/zalpha11 cell line constructing ZC12,749 oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                    AAD22953 standard; DNA; 100
                                                                                                                      Unidentified.
                                                                                                                                                                                                                                                                                                                 26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novak JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 20; Column 149-151; 105pp;
                                                                                                                                                                       inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18;
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V Nelson AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-123547P.
99US-123904P.
99US-142013P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              relates to the isolation of a novel cytokine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sprecher CA,
Dillon SR,
                                                                                                                                                                       sepsis; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hammond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Foster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Holly RD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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03-APR-2001; 2001WO-US10872

WO200177171-A2

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RESULT 14
AAD22954/c
ID AAD229
XX AAD229
XC AAD229
XC AAD229
XX Zalpha
XW Zalpha
KW Inflam
KW Olabet
XX Uniden
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Дb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated soluble zalphall cytokine receptor polypeptide and their cDNA molecules. Zalpha proteins are useful for inhibiting or antagonising the ligand activity-induced proliferation of haematopoietic cells and haematopoietic cell progenitors preferably lymphoid cells which are natural killer cells or cytotoxic T cells. Zalpha is useful for treating immune and inflammatory disorders, for reducing proliferation of neoplastic B or T cells, for suppressing an immune response in a mammal exposed to an antigen or pathogen. Zalpha is useful for treating diseases that require immune regulation including autoimmune diseases such as rheumatoid arthritis, multiple sclerosis, myasthenia gravis, systemic lupus erythematosus (SLE) and diabetes; asthma, ulcerative colitis, inflammatory bowel disease, Crohn's disease, sepsis, viral infection (dengue virus infection) and cancer. The present sequence is an oligonucleotide used for BaF3/KZ134/zalphail cell line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel soluble receptor polypeptides and polynucleotides used as cytokine antagonist for stimulating ligand activity-induced proliferation of hematopoietic cells and for suppressing immune
                                                                       05-APR-2000;
28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                              systemic lupus erythematosus; myasthenia gravis; rheumatoid arthritis; diabetes; autoimmune disease; multiple sclerosis; ulcerative colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Zalphall; cytokine receptor; immunosuppressive; cytostatic;
inflammatory disorder; cell proliferation; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sprecher CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-APR-2000;
                                                                                                                                                  03-APR-2001;
                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                  asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BaF3/KZ134/zalpha11 cell line constructing ZC12,748 oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD22954;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAD22954 standard; DNA; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JUL-2000;
                         (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                   18-OCT-2001
                                                                                                                                                                                                                                                 WO200177171-A2
                                                                                                                                                                                                                                                                                                                                                                         inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OMYZ)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19; Page 213; 243pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                       2000US-194731P.
2000US-222121P.
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2000US-222121P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 33 C; 17 G;
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                                                                                                                                                                                                                                                                                                                                                                         sepsis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 T; 0
                                                                                                                                                                                                                                                                                                                                                                               Crohn's disease;
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76;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                               viral infection;
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                                                                                                                                                                                                               The invention relates to an isolated soluble zalphall cytokine receptor polypeptide and their cDNA molecules. Zalpha proteins are useful for inhibiting or antagonising the ligand activity induced proliferation of haematopoietic cells and haematopoietic cell progenitors preferably lymphoid cells which are natural killer cells or cytotoxic T cells. Zalpha is useful for treating immune and inflammatory disorders, for reducing proliferation of neoplastic B or T cells, for suppressing an immune response in a mammal exposed to an antigen or pathogen. Zalpha is
                                                                                                             myasthenia gravis, systemic lupus erythematosus (SLE) and diabetes; asthma, ulcerative colitis, inflammatory bowel disease, Crohn's disease, sepsis, viral infection (dengue virus infection) and cancer. The present sequence is an oligonucleotide used for BaF3/KZ134/zalpha11 cell line
                                                                                                                                                                                                                                                                                                                                                                                                                       Novel soluble receptor polypeptides and polynucleotides used as cytokine antagonist for stimulating ligand activity-induced proliferation of hematopoletic cells and for suppressing immune
                                                               Sequence 100 BP;
                                                                                                   construction.
                                                                                                                                                                                    useful for treating diseases that require immune regulation including autoimmune diseases such as rheumatoid arthritis, multiple sclerosis,
                                                                                                                                                                                                                                                                                                                                                                         Example 19; Page 213;
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               Local
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18;
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              85.7%;
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Pred. No.
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Mismatches
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Indi RESULT 15 Cytochrome bcl; 07-OCT-2002 ABK88204; ABK88204 standard; DNA; 33 hrome bcl; primer; ss; core protein III2. immunodeficiency virus; human; PCR. cytochrome bcl core protein cDNA specific PCR primer ა 8 AGATTTCTAGGAATTCAA AGATTTCTAGGAATTCAA 18 (first entry) 21 21; cancer; HIV; #2

CN1340524-A.

31-AUG-2000; 2000CN-0119831.

31-AUG-2000; 2000CN-0119831.

20-MAR-2002

(BODE-) BODE GENE DEV CO LID SHANGHAI

Xie

WPI; 2002-436418/47

Polypeptide-human cytochrome polynucleotide for coding it bc1 compound core protein ΙΙ 12. . 21 and

Example 4; Page 19 (disclosure); 34pp; Chinese

This invention relates ţ the cDNA and protein sequences 0f עם novel

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RESULT 16
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one such orrespond to the prevention, diagnosis and treatment by them may be used in the prevention, diagnosis and treatment listed diseases associated with inappropriate expression of the proteins listed diseases associated with inappropriate expression of the proteins listed diseases associated with inappropriate expression of the proteins listed diseases.
                                           proteins, cytochromes, kinesins, cytokines, interferons, interleukins, 6-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encode by them may be used in the prevention, diagnosis and treatment of
                                                                                                            The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                               Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therap; vaccine; amylase; can amyloid protein; angiopoletin; apoptosis related protein; catherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f; complement related protein; cytochrome; kinesin; cytokine; interferinterleukin; G-protein coupled receptor; thioesterase; inflammation multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide-human cytochrome bcl compounds core protein III2. 21. The invention also comprises a method for producing the protein by recombinant DNA technology and a method for the application of the polypeptide in treating diseases such as cancer, HIV infection, etc. The invention also discloses an antagonist against this polypeptide and its therapeutic action, and the application of the polynucleotide to coding this new human cytochrome bcl compound core protein III2.21. The present sequence represents a PCR primer used to clone the human cytochrome bcl compound of the invention.
                                                                                                                                                                                                                                                                                                                                    Shimkets
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27-DEC-2000;
                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                              oligonucleotide
                                                                                                                                                                                                                                                                                                                                                  at low doses. The value especially in cows,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a vaccine against foot and mouth disa (FMD) comprising (in addition to a veterinary vehicle or excipient) an antigen consisting of empty FMD virus capsids, obtained by expression enhanced to the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentaryotic cells of the cDNA of the following regions of the FMD genomentary the control of 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            systemic lupus
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        N
                                                                                l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      potent effective vaccines against foot-and-mouth disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTTCTAGGAAGTCAAA
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                                                                                                                                                                                                                                     A; 7 C; 6 G;
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94.1%;
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ised in an exam
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                                                                                                                    14.8;
No. 2.
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                                                                                                                    .1e+03;
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                                                                                                                                                     Length
                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                  invention.
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RESULT 19
ABL53037/c
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ABL53041/c
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                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                               The present invention relates to a vaccine against foot and mouth disease (FMD) comprising (in addition to a veterinary vehicle or excipient) an antigen consisting of empty FMD virus capsids, obtained by expression in eukaryotic cells of the cDNA of the following regions of the FMD genome: the P1 region encoding the capsid and the region encoding the 3C protease. The vaccine is effective, reliable and stable, and is effective at low doses. The vaccine is useful against foot and mouth disease, especially in cows, sheep, pigs or goats. The present sequence is an oligonucleotide which was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     King
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Virucide; vac
3C protease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide JCA 327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL53041 standard; DNA; 41
                                                                                                                                                                                                                                                                                                                            Sequence 41 BP; 13 A; 6 C; 6 G; 16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Foot and mouth disease virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-2002
                                                                 3C protease;
                                                                             Virucide; vaccine; foot and mouth disease; P1 region; capsid;
                                                                                                     Oligonucleotide JCA
                                                                                                                            29-MAY-2002
                                                                                                                                                                           ABL53037 standard; DNA; 44 BP
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                    WO200200251-A1
                                             Foot and mouth
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                                                                                                                                                                                                                                                        2 GATTTCTAGGAATTCAAA 19
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                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burman
                                                                                                                                                                                                                                   GATTTTTATGAATTCAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            recombinantly produced
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                                             disease virus
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88.9%;
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                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                     Score 14.8;
                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            empty virus capsids as antigens
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                                                                                                                                                                                                                                                                                                     DB 24;
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                                                                                                                                                                                                                                                                                                     Length 41;
                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                0
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RESULT 20
ABL53043/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antigen consisting of empty FMD virus capsids, obtained by expression in eukaryotic cells of the cDNA of the following regions of the FMD genome: the P1 region encoding the capsid and the region encoding the 3C protease. The vaccine is effective, reliable and stable, and is effective at low doses. The vaccine is useful against foot and mouth disease, especially in cows, sheep, pigs or goats. The present sequence is an oligonucleotide which was used in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        King
                                                                                                                                                                                                                                                                       Virucide; vaccine; 3C protease; ds.
                                                                                                                                                           29-JUN-2000; 2000FR-0008437
                                                                                                                                                                                27-JUN-2001; 2001WO-FR02042.
                                                                                                                                                                                                                           WO200200251-A1.
                                                                                                                                                                                                                                                   Foot and mouth disease virus
                                                                                                                                                                                                                                                                                                        Oligonucleotide JCA 329
                                                                                                                                                                                                                                                                                                                               29-MAY-2002
                                                                                                                                                                                                                                                                                                                                                     ABL53043;
                                                                                                                                                                                                                                                                                                                                                                           ABL53043 standard; DNA; 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 44 BP; 13 A; 7 C; 7 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a vaccine against foot and mouth disease (FMD) comprising (in addition to a veterinary vehicle or excipient) an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 26; 79pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2001; 2001WO-FR02042
                                                                                         WPI; 2002-130837/17
                                                                                                                                                                                                      03-JAN-2002
                                                                                                                                     (MERI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           potent effective vaccines against foot-and-mouth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MERIAL
                                                                  potent effective vaccines
                                                                                                                                     MERIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Burman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                  GATTTTTATGAATTCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GATTTCTAGGAATTCAAA
                                                                                                               Burman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 recombinantly produced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                               (first
                                                                                                              Þ,
                                                                                                                                                                                                                                                                                  foot and mouth disease; P1 region; capsid;
                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Audonnet J,
                                                                                                               Audonnet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70.5%;
88.9%;
                                                       produced
                                                                                                                                                                                                                                                                                                                                                                            ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                   N
                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 24
Pred. No. 2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lombard M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 empty virus capsids as antigens
                                                                                                                Lombard
                                                       against foot-and-mouth disease, empty virus capsids as antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and is effective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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0

The present invention relates to a vaccine against foot  $\epsilon$  (FMD) comprising (in addition to a veterinary vehicle or

and mouth dr excipient)

disease

Example 7; Page 30; 79pp; French

recombinantly

empty

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RESULT 21
AAZ23234/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                  The invention provides nucleic acid molecules encoding hepatitis C virus (HCV) non-structural 5B (NS5B) proteins. The HCV NS5B protein can be used in assays to determine antagonistic or agonistic activity of test compounds against HCV. HCV can be detected in biological samples by amplification of the NS5B coding sequence and detection using an oligonuclectide probe (derived from the NS5B nucleotide sequence). The HCV NS5B protein is a viral antigen and can be used in raising an immune response in a mammalian subject. Cell lines comprising the HCV NS5B nucleic acid sequence can be used to assess the functionality of the protein and for assaying test compounds for antagonistic or agonistic
activity. The HCV NSSB protein and nucleic acid sequences are useful in research, diagnostic, therapeutic and pharmaceutical applications, and for use in assays for the identification of efficacious antivixal agents. Sequences AAZ23232-235 represent PCR primers for cloning the HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus; HCV; non-structural 5B; viral antigen; immune response; diagnostic; therapeutic; pharmaceutical;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antigen consisting of empty FMD virus capsids, obtained by expression eukaryotic cells of the cDNA of the following regions of the FMD genom the P1 region encoding the capsid and the region encoding the 3C protease. The vaccine is effective, reliable and stable, and is effect at low doses. The vaccine is useful against foot and mouth disease, especially in cows, sheep, pigs or goats. The present sequence is an especially in cows, sheep, pigs or goats. The present sequence is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9951781-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ23234 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 44 BP; 12 A; 10 C; 9 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 47; 129pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel protein and polynucleotides used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-0CT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HCV NS5B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ23234;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligonucleotide which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1999-620215/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VIROPHARMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GATTTTTATGAATTCAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GATTTCTAGGAATTCAAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                treatments for Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amplifying nested
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0080509.
98US-0090356.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US07404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            70.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NS5B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiviral;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                is effective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FMD genome:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
  HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Sequence 46

BP;

17 A

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14

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0 other

RESULT 23
AAX80718
ID AAX80

AAX80718 standard; DNA; 58

ВP

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RESULT 22
AAV59344
                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                 polypeptide zsig10. zsig10 is involved in mucuos more and the such as adhesion. The products of the invention can be used in the study and treatment of e.g. tumour metastasts, bacterial colonisation, susceptibility to and persistence of infection, microbial infections, AIDS, cystic fibrosis, chronic obstructive pulmonary disease, asthma, AIDS, cystic fibrosis, chronic obstructive pulmonary disease, bronchiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss; human; mucous-mediated function; adhesion; tumour metastasis; bacterial colonisation; microbial infection; AIDS; cystic fibrosis; chronic obstructive pulmonary disease; asthma; Crohn's disease;
                                                                                                                                    Sequence
                                                                                                                                                                  diagnosis
                                                                                                                                                                                                                                                                                                                                          Example 7; Page 94; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                products
                                                                                                                                                                                                                                                                                                                                                                                                              New isolated mucous-associated polypeptide, zsig10 - used to develop
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-531566/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sheppard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9841627-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 zsigl0 primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV59344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV59344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sinonasal inflammatory disease; inflammatory bowel disease; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                      Crohn's
                                                                                                                                                                                                                                                                                      primers AAV59330-V59351 were used in the production of a human ypeptide zsig10. zsig10 is involved in mucous-mediated functions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer;
14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AGATTTCTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                            fibrosis, asthma, bronchitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                  Similarity
                                                                                                                                      58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PO;
                                AGATTTCTAGGAATTCAAATC
                                                                                                                                                                                                                                                                                                                                                                                            for treating e.g. tumour metastasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATATTTATAAGAATTCAGATC
                                                                                                                                                                      and
                                                                                                                                                                                      disease.
                                                                                                                                    BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                      drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZC15,632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0039631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US05251.
                                                                                                                                    A; 19 C; 9 G;
                                                                                                                                                                      screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69.5%;
                                                                                  69.5%;
                                                                                                                                                                                      The products can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.6; DB 20
Pred. No. 2.7e+03;
0; Mismatches 4
                                                                                  Score 14.6;
Pred. No. 2.
                                21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ഗ
34
                                                                   Mismatches
                                                                                                                                    13
                                                                                                                                    Η;
                                                                                                                                                                                                                                                                                                                                                                            or inflammatory bowel disease
                                                                                                                                                                                        also
                                                                                                                                      0
                                                                                    .7e+03;
                                                                                                                                      other;
                                                                                                    DB 19;
                                                                                                                                                                                        be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
                                                                                                                                                                                                                                                                                                                                                                                                microbial
                                                                                                  Length
                                                                                                                                                                                    disease,
for detec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                infections,
                                                                                                     58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46;
                                                                                                                                                                                                      bronchitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                  Gaps
                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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RESULT 24
AAX24706
ID AAX24
XX AAX24
XX AAX24
XX Linke
XX Linke
XX Adipo
KW Adipo
KW Energ
XX Energ
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              systems
tissues
     04-MAR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       helical protein. Zalphal tagged with a Glu-Glu tag at the C-terminus in Pichia methanolica. The Zalphal protein may be useful in the treatment of Fragile-X syndrome and abnormal proliferative disorders e.g. cancer. It can also be used for the growth, differentiation, maintenance and
                                                          WO9910492-A1
                                                                                                            Synthetic
                                                                                                                                                           Adipocyte-specific protein; zsig39; human; fatty acid mergy balance; nutrition; antimicrobial; PCR; primer;
                                                                                                                                                                                                                                            Linker primer ZC14819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   It can also survival of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CEE-Zalphal linker and N-terminal untagged linker were used for construction of a CEE-Zalphal plasmid which expressed the human alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is an oligonucleotide which was used in P for synthesis of C-terminal CEE-Zalphal linker. The C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer; human; alpha helical protein-1; Zalpha1; Fragile-X syndrome; connective tissue; abnormal proliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                    21-JUN-1999
                                                                                                                                                                                                                                                                                                                                                         AAX24706;
                                                                                                                                                                                                                                                                                                                                                                                                        AAX24706 standard; DNA; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalian alpha helical protein-1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-385572/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9929720-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cosmetic improvement; skin tone; elasticity; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer-5 for C-terminal CEE-Zalpha1 linker synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX80718;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ZYMO ) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 AGATTTCTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         58 BP; 17 A; 19 C; 9 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGAATACTAGGAATTCTACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           connective tissues, particularly cardiovascular and epidermal in imparting cosmetic improvements to normal connective has enhancement of skin tone and elasticity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lok S,
                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-0987926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-US26273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69.5%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INC.
                                                                                                                                                                                                                                                                                                                                                                                                              뫔
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.6; DB 20 Pred. No. 2.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>34</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        designated Zalphal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 20;
                                                                                                                                                                                       fatty acid metabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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RESULT 25
AAS00972
ID AAS00
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotides ZC14819, ZC15267 (see AAX24705), ZC14820 (see AAX24707) and ZC15273 (see AAX24704) were used in a PCR amplification to generate an N-terminal untragged zsig39 linker. This was utilised in the construction of a mammalian expression plasmid for zsig39. Zsig39 (see AAW97984) is an adipocyte-specific protein homologue that can be used to modulate fatty acid metabolism (claimed). It may also be used in organ preservation, for cryopreservation, for surgical pretreatment to prevent injury due
                                                           Preparing a double-stranded, circular DNA molecule, involves homologous recombination of one or more donor DNA fragments encoding the protein of interest, with an acceptor plasmid and DNA linkers in host cell -
            The sequence represents a PCR antisense common primer for
                                                                                                                WPI; 2001-256851/26.
                                                                                                                                                                                                                                             27-MAR-2001
                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                      C-terminal
                                                                                                                                                                                                                                                                                                                                                                                       16-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                 AAS00972;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 58 BP; 17 A; 16 C; 9 G; 16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to ischaemia and/or inflammation, and as an antimicrobial agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-204665/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Humes JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-AUG-1998;
                                    Example 1; Columns 25-26; 23pp; English
                                                                                                                                          Raymond CK;
                                                                                                                                                                                             16-OCT-1997;
                                                                                                                                                                                                                    15-OCT-1998;
                                                                                                                                                                                                                                                                       US6207442-B1.
                                                                                                                                                                                                                                                                                                                       homologous recombination; 15632; ss.
                                                                                                                                                                                                                                                                                                                                                              PCR antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAS00972 standard; DNA; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 8; 128; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                   (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein - used to modulate fatty acid metabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGAATACTAGGAATTCTATTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGATTTCTAGGAATTCAAATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sheppard
                                                                                                                                                                                                                                                                                                                                    linker; plasmid construction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                            common primer for C-terminal linker, no.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0056983
                                                                                                                                                                                             97US-0062061
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                                                                                                                                                                                                                     98US-0173043.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      69.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.6; DB 20
Pred. No. 2.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                    PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58;
             a C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0;

linker. The method of the invention comprises preparing

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RESULT 26
ABN83120
                       Query Match
Best Local S
Matches 17
instability, also for cosmetic improvements of the specific antibodies and anti-idiotypic antibodies, for tre proliferation, e.g. cancer, and degeneration. The present represents a PCR primer used in the invention to generate
                                               treating blood vessel diseases, macroorchidism, skin disorders and joint instability, also for cosmetic improvement of the skin or also its specific antibodies and anti-idiotypic antibodies, for treating abnormal
                                                                                                                             has vasotropic, dermatological, cytostatic, and antidegenerative activity. The polynucleotide is used for expressing the protein, for detecting mutations and alterations in transcription of the corresponding gene in fragile-X syndrome patients, and in gene therapy. The polypeptide or its (ant)agonists, are used for treating some of the phenotypic traits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                double-stranded, circular DNA molecule, comprising combining donor DNA fragments encoding the protesin of interest with an acceptor plasmid, and two DNA linkers in a Saccharomyces cerevisiae host cell. The encoding DNA is linked to the acceptor plasmid by homologous recombination of with the linkers and acceptor plasmid to form the closed, circular plasmid. The obtained plasmid is useful for transforming host cells and producing proteins of interest. The method allows for production of a standardised plasmid into which a variety of DNA sequences can be readily inserted and subsequently expressed.
                                                                                             gene in fragile-X syndrome patients, and in gene therapy. The pol or its (ant)agonists, are used for treating some of the phenotypi associated with fragile-X syndrome, for promoting growth, differentiation, maintenance and survival of connective tissues,
                                                                                                                                                                                                              The invention relates to a novel isolated polynucleotide encoding a human alpha helical polypeptide. The polynucleotide of the invention
                                                                                                                                                                                                                                                                                                               New polynucleotide treating phenotypic
                                                                                                                                                                                                                                                                                                                                                                                                Ьķ
                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-DEC-1997;
10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; antidegenerative;
                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-471089/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2002058801-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              degeneration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABN83120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                 OMYZ)
                                                                                                                                                                                                                                                                                                                                                                                                ß
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; alpha helical protein-1; Zalpha1; vasotropic; dermatological;
tatic; antidegenerative; fragile-X syndrome; gene therapy; canc
vessel disease; macroorchidism; skin disorder; joint instabili
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          alpha helical protein-1 C-terminal Zalphal-CBE linker PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14
                                                                                                                                                                                                                                                                 6; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                 ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                             nucleotide encoding human Zalpha1 protein, useful e.g. phenotypic traits of fragile-X syndrome, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGATTTCTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGAATACTAGGAATTCTACTC 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            58 BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             helical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001US-0932679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-067779P
98US-0209525
                                                                                                                                                                                                                                                                22; 27pp; English
                                                                                                                                                                                                                                                                                                                                                                                               DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 19 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       69.5%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                                               Parrish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                               표
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ଫୁ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.6;
                    invention to generate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Τ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          syndrome; gen
kin disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           58;
                                 sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                    C-terminal
                                                                                                                                                                                                                                                                                                                              for
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                                                                              .g. fo
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В S

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RESULT 27
ABK12090
ID ABK1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 17
                                            Matches
                                                     Query Match
Best Local
                                                                                                         its portion that encodes a polypeptide. The polypeptide of the invent comprises 146 or 126 amino acids, also fully defined in the specification. The polypucleotide of the invention is useful for providing an encoded polypeptide that may be used for treating condit associated with abnormal physiology or development e.g. abnormal proliferation (e.g. cancerous conditions) or degenerative conditions. The present nucleic acid sequence represents the C-terminal-CEE-Zalph tagged linker oligonucleotide #2 sequence of the invention. Zalphal is a mammalian alpha helical protein-1.
                                                                                      Sequence
                                                                                                                                                                                                                          The
                                                                                                                                                                                                                                                                     development, degenerative
                                                                                                                                                                                                                                                                                                       New
                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                   K
L
Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         abnormal proliferation; degenerative condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zalpha1-CE linker, for the production of a plasmid human alpha helical protein-1 (Zalpha1) gene.
                                                                                                                                                                                                                                              Example
                                                                                                                                                                                                                                                                                            for
                                                                                                                                                                                                                                                                                                                                                                                              10-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                    10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6303770-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zalpha1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C-terminal-CEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK12090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK12090
                                                                                                                                                                                                                                                                                                                                                                        (OMYZ)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27
                                                                                                                                                                                                            present invention relates to a new polynucleotide sequence, Zalphal, ch comprises 907 nucleotides, fully defined in the specification, or
                                                                                                                                                                                                                                                                   treating conditions associated slopment, e.g. proliferative or snerative conditions
                                                                                                                                                                                                                                                                              polynucleotides encoding mammalian alpha helical protein-1, treating conditions associated with abnormal physiology or elopment, e.g. proliferative or cancerous conditions, or
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 14
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                                                                                                                                                                                                                                              6; Column 39;
                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                  Conklin DC,
                                                                                                                                                                                                                                                                                                                                                                        ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                       83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58
                     AGATTTCTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                alpha helical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AGATTTCTAGGAATTCAAATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGAATACTAGGAATTCTACTC
                                           Conservative
                                                                                      BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                              97US-067779P
                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0209525.
                                                                                      Α;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 19 C;
                                                     69.5%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                                        INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69.5%;
81.0%;
                                                                                                                                                                                                                                                                                                                                                  Parrish
                                                                                       19
                                                                                                                                                                                                                                            25pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein-1; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tagged linker oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancerous condition; to C-terminal~CEE-Zalphal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58
                                                                                      Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9 G;
                                           0;
                                                                                       9
                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                     Score 14.6; DB Pred. No. 2.7e+(
                                                                                                                                                                                                                                                                                                                                                  JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6;
Pred. No. 2.
                                                                                      <u>ن</u>
 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                       13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13 T;
                                                                                      H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7e+03
                                                       .7e+03
                                                                                      other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 В
                                                                24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tagged
                                                                                                                      C-terminal-CEE-Zalphal invention. Zalphal is
                                           Indels
                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linker
                                                                1 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58;
                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the
                                                                                                                                                                                                    invention
                                                                                                                                                                   conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                          Gaps
                                          0;
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RESULT 28
ABV72202/c
ID ABV722
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AAI64395/c
ID AAI64395 standard;
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                                                                                                                                                                                                                                                                                      Query Match
Best Local S
                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                              The specification describes a method for producing a DNA fingerprint of an individual. The method comprises generating restriction fragments to which an adaptor is ligated, amplifying fragments containing a miniature inverted repeat transposable element (MITE) and resolving the amplified fragments. The presence of a certain amplified fragment is correlated to a phenotype. The method is used to characterize the DNA of an individual, to detect polymorphisms, to correlate presence of an amplified fragment with phenotype and to correlate presence of an amplified fragment with phenotype and to correlate presence of molecular manufactures.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Producing a DNA fingerprint of an individual by amplifying fragments containing a miniature inverted repeat transposable element is usefuto detect polymorphisms and correlate genotype with phenotype
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA fingerprint; phenotype; polymorphism; MITE; molecular marker; miniature inverted repeat transposable element; PCR; primer; ss.
            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to detect polymorphisms and particularly in maize -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-SEP-2000; 2000US-0662402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6420117-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABV72202;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV72202 standard;
                                   cell organelle
                                                  Green fluorescence protein; GFP;
                                                                                                                                AAI64395;
                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 5 C; 3
                                                                                                                                                                                                                                                                                                                                                       markers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Column 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-654638/70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wessler SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-SEP-1999;
                                                                           PCR primer #2.
                                                                                                    23-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYGE-) UNIV GEORGIA RES FOUND
                                                                                                                                                                                                                          19
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                                                                                                                                                                                                                                                                            16;
                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                         The
                                                                                                                                                                                                                          ATTTCGAGGAATTGGAATC 1
                                                                                                                                                                                                                                                  ATTTCTAGGAATTCAAATC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                       present PCR primer is used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Casa AM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-153812P
                                      formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of a PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                    entry)
                                                                                                                                                                                                                                                                                      67.6%;
84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                          27 BP
                                       ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                             G; 6 T; 0 other;
                                                                                                                                                                                                                                                                        Score 14.2; DB 4.,
Pred. No. 3.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INC.
                                                  drug
                                                  screening; PCR primer;
                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is useful
                                                                                                                                                                                                                                                                            Gaps
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RESULT 30
ANI64405/c
ID ANI644
XX ANI644
XX ANI644
XX Green
DT 23-NOV
XX Green
KW Green
XX Uniden
XX Uniden
XX Uniden
XX UP2001
XX UP1; 2
XX UP1; 
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a trap vector containing, from 5' - to 3'-, an intron and splice acceptor sequence, a DNA sequence encoding green fluorescence protein (GFP) in which methionine and valine at the N-terminal are deleted, an internal ribosome binding site (IRES), a DNA sequence encoding a selective marker, and polyadenylated signal sequence. The invention can be used in the screening of drugs inhibiting or promoting the formation of cell organelles. The present sequence is a PCR
The present invention relates to a trap vector containing, from 5'- to 3'-, an intron and splice acceptor sequence, a DNA sequence encoding green fluorescence protein (GFP) in which methionine and valine at the N-terminal are deleted, an internal ribosome binding site (IRES), a DNA sequence encoding a selective marker, and polyadenylated signal sequence. The invention can be used in the screening of drugs inhibiting or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP2001157588-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene trap using a green fluorescence protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUN-2001.
                                                                                                                                                                                                                                                                          Gene trap using a green
                                                                                                                                                                                                                                                                                                                                    WPI; 2001-592442/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell organelle formation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Green fluorescence protein; GFP; drug screening; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI64405;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI64405 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 27 BP; 6 A; 6 C; 7 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer, which was used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 6; 17pp; Japanese.
                                                                                                                                                                                                                Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP2001157588-A.
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                                                                                                                                                                                                                                                                                                                                                                                                 (OSAB-) ZH OSAKA BIOSCIENCE KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19 AGCTAGCTAGGAATTCAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGATTTCTAGGAATTCAAA 19
                                                                                                                                                                                                                Page 7; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99JP-0343222
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84.2%;
                                                                                                                                                                                                                                                                                fluorescence protein
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Pred. No. 4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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G; Conservative (
AGATTTCTAGGAATTCAAA

Matches

16;

Best Local Similarity

84.2%;

0; 19

Pred. No. 4.1.
); Mismatches

. 4.1e+03; 3;

Indels

0;

Gaps

0;

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AAI76494/c
ID AAI76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 16
                                                                                                            sequences (I), which contain single nucleotide polymorphisms (SNPs).

AAM53314 to AAM53329 represent peptides related to human polymorphic polymucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations or deletions in a patient's genome that affect the activity of polypeptides by expressing inactive proteins or to supplement the patients own production of polypeptide. Additionally, (I) and its complementary sequences may also be used as DNA probes in diagnostic assays to detect and quantitate the presence of similar nucleic acids in semantace and therefore which better may be in mad of restractive.
                                       assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. The polypeptides encoded by (1) may be used as antigens in the production of antibodies specific for polymorphic polypeptides. The antibodies may also be used to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the
 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; genome; gene ther protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                               AAI73060 to AAI79867 represent isolated human polymorphic
                                                                                                                                                                                                                                                                                                                                                                        Polymorphic nucleic acid sequences, useful in genetic testing and
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-356160/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-NOV-2000; 2000US-0726173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human silent SNP containing nucleic acid SEQ:3435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoting the form primer, which was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19
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                                                                                                                                                                                                                                                                                                                               Page 1102; 2653pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGCTAGCTAGGAATTCAAA 1
BP;
                             polymorphic polypeptides
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84.2%;
10 C; 3 G;
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21 T; 0
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                          samples
other;
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                                                                                                                                                                                                                                                                                                  polynucleotide
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Query Match

67.6%;

Score 14.2;

DB 22;

Length

51;

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114-MAY-1999;
18-MAY-1999;
19-MAY-1999;
21-MAY-1999;
24-MAY-1999;
25-MAY-1999;
27-MAY-1999;
27-MAY-1999;
27-MAY-1999;
01-JUN-1999;
01-JUN-1999;
01-JUN-1999;
01-JUN-1999;
01-JUN-1999;
01-JUN-1999;
01-JUN-1999;
                                                                                                                                                                      30-APR-1999;
04-MAY-1999;
05-MAY-1999;
06-MAY-1999;
06-MAY-1999;
07-MAY-1999;
11-MAY-1999;
                                                                                                                                                                                                                                  25-MAR-1999
29-MAR-1999
01-APR-1999
06-APR-1999
16-APR-1999
16-APR-1999
19-APR-1999
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23-APR-1999
23-APR-1999
28-APR-1999
30-APR-1999
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05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybridisation assay; genetic mapping; gene expression protein identification; signal transduction pathway;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana.
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-1999;
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                                                                                                                                                                                                                                                                                                                                                                                   2000EP-0301439
99US-0136392.
99US-0136782.
99US-0137222.
99US-0137528.
99US-0137528.
99US-0137502.
99US-0138094.
99US-0138647.
99US-0139119.
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99US-0135124.
99US-0135353.
99US-0135629.
99US-0136021.
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99US-0132486.
99US-0132487.
99US-0132487.
99US-0134256.
99US-0134218.
99US-0134218.
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99US-0131449.
99US-0132048.
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99US-0134370.
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99US-0130510
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                                                           Query Match
Best Local S
Matches 16
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20-AUG-1999;
20-AUG-1999;
20-AUG-1999;
23-AUG-1999;
25-AUG-1999;
27-AUG-1999;
27-AUG-1999;
31-AUG-1999;
31-AU
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 36
                                                           16; Conservative
 AGATTTCAAGGAACACAAA
                      AGATTTCTAGGAATTCAAA
                                                                                                                       99US-0148341
99US-0148684
99US-0149368
99US-0149426
99US-0149426
99US-0149922
99US-0151066
99US-01510884
99US-01510880
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99US-0169231
99US-0169231
99US-016081
99US-016081
99US-016081
99US-0161359
99US-0161361
                                                                             67.6%;
84.2%;
                                                          Score 14.2; DB 21;
Pred. No. 4.1e+03;
0; Mismatches 3;
   54
                                 19
                                                                                           Length
                                                                  Indels
                                                                                                61;
                                                                  0;
                                                                  Gaps
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16-JUN-1999
17-JUN-1999
18-JUN-1999
19-JUN-1999
19-JUN

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3-0139763 3-0139819 3-0149353 3-0140353 3-0140695 3-0140823 3-0141287 3-0141287 3-0141287

3-0139454 3-0139455 3-0139457 3-0139459 3-0139459 3-0139460 3-0139462 3-0139463 3-0139463

S-0139452. S-0139453. S-0139492.

1999; 1999;

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1999; 1999; 1999; 1999;

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3-0142055.
3-0142803.
3-0142803.
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3-0142977.
3-0144065.
3-0144065.
3-014433.
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3-014508.
3-014708.
3-0147192.
3-0147193.
3-0148171.
3-0148171.

1999;

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RESULT 34
ABN41529/c
ID ABN415
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AC ABN415
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AC ABN415
XX
DT 15-JUL
XX
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AAT86363/c
                                                                                                                                                         Matches
                                                                                                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                       This is the sequence of probe F-alpha derived from the sequence of the non-transcribed intergene region (IGR) FI from Candida krusei (AAT86361). The sequence is derived from a ribosomal cistron located in the region between the 25S and 18S ribosomal DNA genes. Probes (see AAT86362-T86365) derived from the IGR can be used for specific and/or intraspecific detection, identification and/or differentiation of yeast for medical diagnosis or for monitoring industrial processes, specifically for Candida krusei and Geotrichum candidum.
            15-JUL-2002
                                                        ABN41529
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                            Claim 7; Page 30; 49pp;
                                                                                                                                                                                                                                                                                                                                                            process control - comprises fragment derived from ribosomal the non-transcribed intergene region that contains repeated
                                                                                                                                                                                                                                                                                                                                                                        Probe for detection and characterisation of yeast for diagnosis process control - comprises fragment derived from ribosomal DNA
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-489661/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-MAR-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C. krusei non-transcribed intergene region-derived probe
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                                                                                                                                                                    Similarity
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                                                      standard;
                                                                                                                                  AGATTTCTAGGAATTCAAA 19
                                                                                                             AGATTTCAAAAATTCAAA
                                                                                                                                                                                                     BP;
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                                                                                                                                                         Conservative
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           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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2..7
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                                                        DNA;
                                                                                                                                                                   67.6%;
84.2%;
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                                                         60
                                                                                                                                                                                                                                                                                                                            French.
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                                                                                                                                                         0,
                                                                                                               47
                                                                                                                                                                                                    G; 48 T; 0 other;
                                                                                                                                                                             Score 14.2; DB 18;
                                                                                                                                                                   Pred. No. 4.1e+03;
                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IGR; ribosome; cistron;
                                                                                                                                                         Indels
                                                                                                                                                                           Length
                                                                                                                                                                              81;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25S; 18S;
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20-MAY-1997 AAT58221;

(first entry)

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AAT58221/
ID AAT5
XX
AC AAT5
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AC AAT5
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XX
DT 20-M
XX
                                                                                                                                                                         RESULT 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC biological sample, in expression profiling studies, in qualitatively or CC quantitatively characterising the corresponding transcriptome, and in CC detecting RNA transcripts and splice variants of human or animal CC transcriptomes. The libraries may also be used as specialised mini CC particular biological or pathological state, and so allowing the CC particular biological or pathological state, and so allowing the CC detection of tissue—and pathology-specific genes such as those genes CC only expressed in specific tissue under a specific pathological CC condition; to detect developmental specific genes, and to detect RNA CC transcripts and splice variants of a transcriptome of a patient suffering CC from a particular disorder. ABN27253 to ABN59589 represent on incompletide sequences from rats, humans and mice, which are used in the exemplification of the present invention.

CC N.B. The sequence data for this patent did not form part of the printed consisting the control of the present invention.
                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multipl transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to set of messenger RNAs transcribed from a given transcription unit of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue, pathology, and devolopmental forderesting tissue, pathology, and
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                        AAT58221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   splice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human spliced transcript detection oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (COMP-) COMPUGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-JUL-2001; 2001WO-IB01903
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                 52
                                                                                                                                                                                                                                                                                               N
                                                                                                                                                                                                                                                                                                                                                14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mouse; rat; splice transcript; detection; RNA transcript;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   variant; transcriptome; oligonucleotide library; ss
                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                        standard;
                                                                                                                                                                                                                                                                                             GATTTCTAGGAATT
                                                                                                                                                                                                                                              GATTTCTAGGAATT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID 14277; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wasserman
                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes oligonucleotide libraries for
                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-221607P
2001US-287724P
                                                                                                                                                                                                                                                                                                                                                                                                                                            17 A; 12 C;
                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                      66.7%; but
100.0%; Pr
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                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                            15
                                                                                                                                                                                                                                                                                                                                                                     Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 16 T;
                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                     DB 24;
5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0 other;
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                                                                                                                                                                                                                                                                                                                                                                                         , 24;
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                                                                                                                                                                                                                                                                                                                                                                                            Length 60;
                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IJ
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RESULT 36
AAV04870/c
ID AAV048
XX AAV048
XX 30-APR
DT 30-APR
XX PCR px
XX Glomer
KW Fibror
KW PTPase
KW PCR pr
XX
OS Synthe
XX
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                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Six Candida genes have been isolated, which encode an apparent CDC25 phosphatase (TYP1), a p13suc1 homolog (CKS1), a cyclin dependent kinase (CDK1), a cyclin (CYB1), a CDK-activating kinase catalytic subunit (MOC1), and a Map kinase (CMK1) (AAT64446 to AAT644451). The TYP1 polypeptide and nucleic acid is claimed, where TYP1 is at least 75% homologous to the amino acid sequence given in Seq 2, according to the claims of the specification. According to the disclosure, Seq 2 encodes CKS1 (AAT64446) and Seq 1 encodes
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The products may be used in reagents and assays which permit the rapid detection and evaluation of Candida yeast infections and for identifying cpds. which have antifungal properties and which may be used as anti-mycotic agents. Such agents can be used therapeutically, as well as, for example, preservatives in foodstuff, feed supplement for promoting weight gain in livestock, or in disinfectant formulations for treatment of non-living matter, e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cell-cycle regulatory proteins from Candida - useful as anti-mycotic agents, as a preservative or in reagents and assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cyclin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 7; Page 37; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-043149/04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cottarel G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9639527-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Candida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYP1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Candida
                                                           Glomerular epithelial protein 1; GLEPP1; podocyte; gene therapy; fibronectin type III-like repeat; protein tyrosine phosphatase do pripase domain; glomerular disorder; low level; GLEPP1 expression;
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                  decontaminating hospital equiment and rooms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MITO-) MITOTIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-DEC-1996
                                                                                                                                                        30-APR-1998
                                                                                                                                                                                        AAV04870
                                                                                                                                                                                                                     AAV04870 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYP1 (AAT64447).
                                                PCR primer;
                                                                                                                         PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CKS1; CDK1; CYB1; MOC1; CMK1; cell-cycle regulatory protein;
da; anti-mycotic; antifungal; preservative; yeast;
n; kinase; phosphatase; ss.
                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                           3 ATTTCTAGGAATTCAAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CYB1
                                                                                                                                                                                                                                                                                                  ATTACTATGAATTCAAA 4
                                                                                                                         2 used to amplify the PTPase domain of GLEPP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Damagnez V,
                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 10 A; 6 C; 4 G; 10 T; 0 other;
                                                   SS
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene
                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96WO-US08807.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer.
                                                                                                                                                                                                                                                                                                                                                                          65.7%;
88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Draetta
                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                        Score
                                                                                                                                                                                                                                                                                                                                                                            Pred.
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                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                             DB 18;
                                                                                                                                                                                                                                                                                                                                                             2;
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                                                                                 domain;
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Synthetic

New Candida cyclin activated kinase and screening for its inhibitors -

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for

generating vaccines

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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein 1 (GIEPPI) protein tyrosine phosphattase (PTPase) segment. The PCR product was cloned to produce a fusion protein, which was tested to confirm that the PTPase domain of GIEPPI had PTPase activity. The GIEPPI protein is primarily expressed in mammalian podocytes. The protein has a large extracellular domain with a sequence containing a fibronectin type III-like repeats, a hydrophobic transmembrane segment and a single intracellular protein tyrosine phosphatase (PTPase) domain. The GIEPPI nucleic acids are used for gene therapy of glomerular through the protein tyrosine phosphatase (PTPase) domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 {\tt DNA} encoding glomerular epithelial protein 1 - useful for therapy of glomerular disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30 BP; 7 A; 6 C; 4 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primers AAV04869-70 were used to amplify the glomerular epithelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Column 14; 47pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1994;
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                                                                                                                                                                                                                                                                                    07-FEB-2000
                                                                                                                                                                                                                                                                                                         AAZ08279;
                                                                                                                                                                                                                                                                                                                               AAZ08279
                                                                                                                          05-MAY-1999;
                                                                                                                                                11-NOV-1999
                                                                                                                                                                       WO9957536-A2
                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                  CYB1-VP16
                                                                                                                                                                                                                           Cell cycle regulatory protein; CYB1 gene; Candida; coding sequence; EcoRI; restriction site; primer; amplify; subclone; pJG4-5 vector;
                                                                                                                                                                                                                                                          PCR primer-1 used for amplification of Candida CYB1
                                   WPI; 2000-038847/03
                                                                                                     05-MAY-1998;
                                                                               (MITO-) MITOTIX INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                                                                                                                                                                                                             w
                                                                                                                                                                                                                          restriction site; primer; amplify;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                      ATTTCTAGGAATTCAAA 19
                                                                                                                                                                                                                                                                                                                                                                                    AGTCCTAGGAATTCAAA
                                                                                                                                                                                                                  fusion protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wiggins RC;
                                                         Cottarel
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                            99WO-US09878
                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             65.7%;
88.2%;
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                                                         Damagnez
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                                                                                                                                                                                                                                                                                                                                 ВP
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Pred. No. 6.
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                                                          Rudolph
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                                                          'n
                                                                                                                                                                                                                                                                                                                                                                                                                                     2;
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                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                           30;
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RESULT 38
ABN47639
biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition; to detect developmental specific genes; and to detect RNA transcripts and splice variants of a transcriptome of a patient sufferir
                                                                                                                                                                                                                                                                                                                                        The present invention describes oligonucleotide libraries for detecting messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multiple transcription units that populate a genome. The library comprises several oligonucleotides, each capable of hybridising selectively to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developmental-specific genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUL-2000;
02-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUL-2001; 2001WO-IB01903
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Page 60; 109pp; English.
                                                                                                                                                                                                                                                          genome, which encodes one or more messenger RNA splice variants. oligonucleotide libraries are useful for detecting mRNAs from a
                                                                                                                                                                                                                                                                                                             eral oligonucleotides, each capable of hybridising selectively to a of messenger RNAs transcribed from a given transcription unit of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     variant;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID 20387; 47pp;
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anscriptome; oligonucleotide library; ss.
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88.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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Pred. No. 6.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English
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set of messenger RNAs transcribed from a given transcription unit of the genome, which encodes one or more messenger RNA splice variants. The oligonoucleotide libraries are useful for detecting mRNAs from a biological sample, in expression profiling studies, in qualitatively characterising the corresponding transcriptome, and in detecting RNA transcripts and splice variants of human or animal transcriptomes. The libraries may also be used as specialised mini libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue- and pathology-specific genes such as those genes only expressed in specific tissue under a specific pathological condition, to detect developmental specific genes, and to detect RNA transcripts and splice variants of a transcriptome of a patient suffer
                                                                                                                                                                                                                                                                                                                                                                                       messenger RNAs that populate a (sub-)transcriptome, where the (sub-)transcriptome comprises messenger RNAs transcribed from multranscription units that populate a genome. The library comprises
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                                                                                                                                                                                                                                                                                                                                                                      several
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Pred. No. 6
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N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                           Primer #2 corresponds to nucleotides 5'-3618-3639-3'of ApoE. This plus-strand primer is used with minus strand primer #4 (3'-4220-4241-5', see AAQ32874) to amplify exon 4 of the human apolipoprotein gene. The epsilon 2, 4 and 7 mismatch mutations occur in this region, at positions 3883, 3745 and 4141 and 4144, respectively. Three sets of four oligonucleotide probes were prepared to distinguish the wild-type from the mutant base at the mismatch position for both the plus and the minus strands. The probe sets are AAQ32875-6 and AAQ32883-4 for epsilon 2, AAQ32877-8 and AAQ32885-6 for epsilon 4 and AAQ32881-2 and AAQ32889-90 for epsilon 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Testing apolipoprotein E genotype - using polymerase chain reaction primers and labelled allele-specific oligonucleotide probe for hybridisation to amplified deoxyribonucleic acid
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epsilon 2; epsilon 4; epsilon 5; epsilon 7; ss.
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PRIOR FILING DATE: 2001-06-22
NUMBER OF SEQ ID NOS: 230
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 30
SEQ ID NO 30
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                 ; OTHER INFORMATION: Synthetic oligonucleotide US-09-954-987B-30
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APPLICANT: Grayson B. Lipford
APPLICANT: Grayson B. Lipford
APPLICANT: Hermann Wagner
TITLE OF INVENTION: PROCESS FOR HIGH THROUGHPUT SCREENING
TITLE OF INVENTION: COG-BASED IMMUNO-AGONIST/ANTAGONIST
FILLE REFERENCE: C1041/7016 (AWS)
CURRENT APPLICATION NUMBER: US/09/954,987B
CURRENT FILLING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: US 60/233,035
PRIOR FILLING DATE: 2000-09-15
PRIOR FILLING DATE: 2000-09-15
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PRIOR FILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/300,210
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9 US-09-765-272-409
8 US-08-929-464-4
9 US-09-778-320-29
9 US-09-910-689-29
14 US-10-010-742-299
15 US-10-11-877-31
11 US-9-885-453-11
11 US-9-885-453-11
11 US-9-885-453-11
12 US-10-175-247-5
13 US-10-175-247-5
14 US-10-161-547-35
15 US-10-161-547-35
16 US-10-161-547-35
17 US-09-765-272-415
18 US-10-210-152-264
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19 US-10-217-632-175486
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      Score 18;
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      DB 11;
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Sequence 409, App
Sequence 4, Appl
Sequence 299, App
Sequence 299, App
Sequence 31, Appl
Sequence 31, Appl
Sequence 47, Appl
Sequence 5, Appli
Sequence 26, Appli
Sequence 26, Appli
Sequence 27, Appli
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Sequence 5, Appli
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TITLE OF INVENTION: METHOD FOR SCREENING FOR TITLE OF INVENTION: FACTOR ACTIVITY FILE REFERENCE: 26757-704 CURRENT APPLICATION UNMBER: US/09/877,705A CURRENT FILING DATE: 2001-08-16 NUMBER OF SEQ ID NOS: 162 SOFTWARE: PatentIn version 3.1 SEQ ID NO 92 LENGTH: 21
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US-09-877-705A-91
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RESULT 4
US-09-877-738A-91
; Sequence 91, Application US/09877738A
; Publication No. US20030022173A1
; GENERAL INFORMATION:
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; OTHER INFORMATION: Transcription factor probe PP92
US-09-877-705A-92
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CURRENT APPLICATION NUMBER: US/09/877,705A
CURRENT FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 162
SOFTWARE: PatentIn version 3.1
SEQ ID NO 91
LENGTH: 21
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Publication No. US20030008283A1
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TITLE OF INVENTION: METHOD FOR SCREENING FOR DRUG CANDIDATES FOR MODULATING TRANSCRIF
TITLE OF INVENTION: FACTOR ACTIVITY
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ORGANISM: Artificial sequence
FEATURE:
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                                                                                                                                                                                                                                                                               ORGANISM: Artificial sequence
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Matches
                                FILE REFERENCE: 26757-704
CURRENT APPLICATION NUMBER: US/09/877,705A
CURRENT FILING DATE: 2001-08-16
NUMBER OF SEQ ID NOS: 162
SOFTWARE: Patentin version 3.1
SEQ ID NO 154
LENGTH: 63
                                                                                                                                                                                       Publication No. US200
GENERAL INFORMATION:
APPLICANT: Li, Jason
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SEQ ID NO 92
LENGTH: 21
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LENGTH: 21
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CURRENT APPLICATION NUMBER: US/09/877,738A
CURRENT FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 162
                                                                                                                                                     TITLE OF INVENTION: METHOD FOR SCREENING FOR DRUG CANDIDATES FOR MODULATING TRANSCRITITLE OF INVENTION: FACTOR ACTIVITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: METHOD AND KIT FOR ISOLATING DNA PROBES THAT BIND TO ACTIVATED TITLE OF INVENTION: TRANSCRIPTION FACTORS
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TYPE: DNA ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Transcription factor probe PP92
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ORGANISM: Artificial
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Pred. No.
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Matches
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                                                                                      SOFTWARE: Fa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. US20020128446A1
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Best Local
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TITLE OF INVENTION: WETHOD AND KIT FOR ISOLATING TITLE OF INVENTION: TRANSCRIPTION FACTORS
FILE REFERENCE: 26757-701
CURRENT APPLICATION NUMBER: US/09/877,738A
CURRENT FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 162
                                                                                                                                                      PRIOR FILING DATE: EARLIER FILING NUMBER OF SEQ ID NOS: 115
                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/923,246
CURRENT FILING DATE: 2001-08-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hammond, Angela K.
TITLE OF INVENTION: NOVEL CYTOKINE ZALPHALL LIGAND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial sequence
ORGANISM: Artificial Sequence
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                                                            ENGTH:
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Dillon, Stacey R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Johnston, Janet V.
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D. US20030022173A1
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100.0%; Pred. No.
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Pred. No.
                                                                                                                                                                                       DATE: 1999-07-01
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87;
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CTHER INFORMATION: Oligonucleotide ZC12749
US-09-923-246-59
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US-09-923-246-60/c
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 60
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Best Local Similarity
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APPLICANT:
                                                                                                        APPLICANT: Sprecher, Cindy A.
APPLICANT: No. US2002013/677Alak, Julia
APPLICANT: West, James W.
APPLICANT: Presnell, Scott R.
APPLICANT: Holly, Richard D.
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APPLICANT: Presnell, Scott R.
APPLICANT: Sprecher, Cindy A.
APPLICANT: Foster, Donald C.
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/522,217
PRIOR FILING DATE: EARLIER FILING DATE: 2000-03-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/123,904
PRIOR FILING DATE: EARLIER FILING DATE: 1999-03-11
PRIOR FILING DATE: EARLIER EARLIER APPLICATION NUMBER: US 60/142,013
PRIOR FILING DATE: EARLIER FILING DATE: 1999-07-01
PRIOR FILING DATE: EARLIER FILING DATE: 1999-07-01
                                                                    APPLICANT: Presnell, Scott R.
APPLICANT: Holly, Richard D.
APPLICANT: Nelson, Andrew J.
ATTILE OF INVENTION: SOLUBLE ZALPHA11 CYTOKINE RECEPTORS
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                  CURRENT APPLICATION NUMBER: US/09/825,561A
CURRENT FILING DATE: 2000-04-05
                                                        FILE REFERENCE: 00-22
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PRIOR APPLICATION NUMBER: US 60/194,731
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ORGANISM: Artificial Sequence
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18; Conserv
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Dillon, Stacey R.
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100.0%; Pr
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Pred. No.
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Pred. No.
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95;
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                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Oligonucleotide primer ZC12748 US-09-825-561A-49
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 48
LENGTH: 100
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                                                                                                                                                             Sequence 43, Application US/09892949 Publication No. US20030096339A1
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Best Local Similarity
                                                                                                                                                GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 2000-07-28
                                                                                     APPLICANT: Sprecher, Cindy A. APPLICANT: Presnell, Scott R. APPLICANT: Gao, Zeren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 00-22
CURRENT APPLICATION UMBER: US/09/825,561A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/194,731
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
                                    APPLICANT:
APPLICANT:
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APPLICANT: Holly, Richard D.
APPLICANT: Nelson, Andrew J.
TITLE OF INVENTION: SOLUBLE ZALPHA11 CYTOKINE RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sprecher, Cindy A. APPLICANT: No. US20020137677Alak,
                  APPLICANT: Whitmore, Theodore E.
APPLICANT: Kuliper, Joseph L.
APPLICANT: Maurer, Mark F.
TITLE OF INVENTION: CYTOKINE RECEPTOR ZCYTOR17
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SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: West, James W. APPLICANT: Presnell, Scoti
FILE REFERENCE: 00-42
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100.0%;
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Pred. No.
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Pred. No.
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RESULT 14
US-10-282-622-17
J Sequence 17, Application US/10282622
Publication No. US20030134390A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 44
LENGTH: 100
TYPE: DNA
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Best Local Similarity
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APPLICANT: Presnell, Scott R
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                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/267,963
PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: US 60/214,282
PRIOR FILING DATE: 2000-06-26
PRIOR APPLICATION NUMBER: US 60/214,955
PRIOR APPLICATION NUMBER: US 60/214,955
PRIOR FILING DATE: 2000-06-29
DEFIOR FILING DATE: 2000-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Sprecher, Cindy A. APPLICANT: Presnell, Scott R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/892,949
CURRENT FILING DATE: 2001-06-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 00-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: CYTOKINE RECEPTOR ZCYTOR17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/214,955
PRIOR FILING DATE: 2000-06-29
PRIOR APPLICATION NUMBER: US 60/267,963
PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 93
                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Oligonucleotide primer ZC12748
                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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PRIOR APPLICATION NUMBER: US 60/214,282
PRIOR FILING DATE: 2000-06-26
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CURRENT FILING DATE: 2001-06-26
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Pred. No.
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Pred. No.
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FILE REFERENCE: 01-37
CURRENT APPLICATION NUMBER: US/10/282,622
CURRENT FILLING DATE: 2002-10-28
PRIOR APPLICATION NUMBER: 60/337,586
PRIOR APPLICATION NUMBER: 60/337,586
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PASISEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 100
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                                                      US-10-351-157-44
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Sequence 44, Application No. US20 GENERAL INFORMATION:
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Publication No. US20030134390A1
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Best Local Similarity
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APPLICANT: No. US20030134390Alak, Julia E.
TITLE OF INVENTION: ZALPHA11 LIGAND ANTAGONISTS
FILE REFERENCE: 01-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/282,622
CURRENT FILING DATE: 2002-10-28
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APPLICANT: West, Jan
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APPLICANT: No. US20030134390Alak, Julia E
                                                                                                                                                                                                                                                                          OTHER INFORMATION: oligonucleotide primer ZC12,748
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                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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ORGANISM: Artificial Sequence
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18; Conserva
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                                  Application US/10351157
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APPLICANT: Sprecher, Cindy A.
APPLICANT: Gao, Zeren
APPLICANT: Kuijper, Joseph L
APPLICANT: Dasovich, Maria M
APPLICANT: Grant, Francis J.

Kuijper, Joseph L. Dasovich, Maria M. Grant, Francis J. Presnell, Scott R. Whitmore, Theodore

APPLICANT: APPLICANT: APPLICANT:

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APPLICANT: APPLICANT: APPLICANT:

Dillon,

Stacey R.

Gross, Jane A.

US20030215838Alak, Julia

FILE REFERENCE: 02-02

TITLE OF INVENTION: CYTOKINE RECEPTOR ZCYTOR17 MULTIMERS

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                                                                                            ; OTHER INFORMATION: Oligonucleotide primer ZC12748 US-10-351-157-45
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SEQ ID NO 44
LENGTH: 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sprecher, C
APPLICANT: Gao, Zerer
APPLICANT: Kuijper, J
APPLICANT: Dasovich,
APPLICANT: Grant, Fra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local :
                                Matches
                                             Query Match
Best Local Similarity
                                                                                                                                                                                       SEQ ID NO 45
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CURRENT FILING DATE: 2003-01-21
PRIOR APPLICATION NUMBER: US 60/435,361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/350,325 PRIOR FILING DATE: 2002-01-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2002-12-19
                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                     PRIOR FILING DATE: 200 NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/389,108 PRIOR FILING DATE: 2002-06-14
                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/351,157
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                            LENGTH: 100
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/350,325
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                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 02-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                         FEATURE:
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                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/435,361 FILING DATE: 2002-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
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                                18;
 1 AGATTTCTAGGAATTCAA 18
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Dasovich, Maria M.
                                                                                                                                                                                                                                                                                                                                                                                                                         Hammond, Angela K.
No. US20030215838Alak, Julia E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Presnell, Scott R.
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                                Conservative
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                                                                                                                                                                                                                                      2002-01-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Theodore E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              183
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100.0%;
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100.0%;
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Pred. No.
                                0;
                                                Score 18;
Pred. No.
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                                  Mismatches
                                                DB 13;
95;
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                                                                Length 100;
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         APPLICANT: Hammond, Angela K.
TITLE OF INVENTION: NOVEL CYTOKINE ZALPHA11 LI.
FILE REFERENCE: 99-16
CURRENT APPLICATION NUMBER: US/10/295,723
CURRENT FILING DATE: 2002-11-15
PRIOR APPLICATION NUMBER: 09/522,217
PRIOR FILING DATE: 2000-03-09
PRIOR APPLICATION NUMBER: US 60/124 F47
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Best Local Similarity
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APPLICANT:
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NUMBER OF SEQ ID NOS: 115
SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: Presnell, Scott R.
APPLICANT: Sprecher, Cindy A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NOVEL CYTOKINE ZALPHA11 LIGAND
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PRIOR FILING DATE: 1999-03-09
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PRIOR FILING DATE: 1999-03-09
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Dillon, Stacey R.
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                                                                                                                                                                                  Johnston, Janet V.
Nelson, Andrew J.
                                                                                                                                                                                                                          Gross, Jane A.
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                                                                                                                                                                 Nelson, Andrew J.
Dillon, Stacey R.
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                                                                                                                                                                                                                                                               Donald
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100.0%;
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Pred. No. 95;
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Gaps

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RESULT 20
US-10-325-606-1/c
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Query Match
Best Local Similarity
Matches 16; Conserv
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Publication No. US20030216295A1
GENERAL INFORMATION:
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PRIOR TILING DATE: 1999-03-11
PRIOR APPLICATION NUMBER: US 60/142,013
PRIOR FILING DATE: 1999-07-01
NUMBER OF SEQ ID NOS: 115
                                                                                                                                                                           TELEFAX: 619-535-8949 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: P-LJ 1113
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-535-9001
                                                                                          SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/369,754
FILING DATE: 06-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: CAMPBELL, CATHRYN
                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/325,606
FILING DATE: 20-Dec-2002
                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: CAMPBELL AND FLORES
STREET: 4370 LA JOLLA VILLAGE DRIVE, STE. 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: E-SELECTIN-DEPENDENT CELL ADHESION ANTAGONISTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FUKUDA, MINORU
SAWADA, RITSUKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    38 AGATTTCTAGGAATTCAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
                                                                                                            TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                 TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: CALIFORNIA COUNTRY: UNITED STATES
                                                                                                                                                          LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
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                        73.3%;
94.1%;
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100.0%;
        0; Mismatches
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                        Score 15.4;
Pred. No. 1
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Pred. No.
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                        1.2e+03;
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                                      DB 13;
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                                     Length
       Indels
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   Gaps
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Query Match
Best Local Similarity
Thes 16; Conserve
                                                                                                  ; OTHER INFORMATION: oligonucleotide JCA327 used
; OTHER INFORMATION: a fragment of about 3460 bp
US-10-327-481A-23
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                                                                                                                                                                                                                                            SEQ ID NO 23
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Matches
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LENGTH: 41
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                                                                                                                                                                                                                                                                                                                         APPLICANT: Audonnet, Jean-Christophe F.
APPLICANT: Lombard, Michel F.A.
TITLE OF INVENTION: Vaccine against Foot-and-Mouth Disease
FILE REFERENCE: 454313-3178
CURRENT APPLICATION NUMBER: US/10/327,481A
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: PCT/FR01/02042
PRIOR APPLICATION NUMBER: FR 0008437
PRIOR APPLICATION NUMBER: FR 0008437
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Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: King, Andrew M.Q. APPLICANT: Burman, Alison J APPLICANT: Audonnet, Jean-C
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APPLICANT: Lombard, Michel F.A.
TITLE OF INVENTION: Vaccine against Foot-and-Mouth Disease
FILE REFERENCE: 454313-3178
CURRENT APPLICATION NUMBER: US/10/327,481A
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: PCT/FR01/02042
PRIOR FILING DATE: 2001-06-27
PRIOR FILING DATE: 2001-06-27
                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Ver.
                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 41
                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: FR 0008437
PRIOR FILING DATE: 2000-06-29
NUMBER OF SEQ ID NOS: 41
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                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial sequence
                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: oligonucleotide JCA325 used OTHER INFORMATION: a fragment of about 3470 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial sequence
                                                                                                                                                                                                                           ENGTH: 41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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No. US20040001864A1
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                                  70.5%;
88.9%;
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                                    Score 14.8; DB 12;
Pred. No. 2.6e+03;
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Pred. No. 2.6e+03;
0; Mismatches 2
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                                                                                                                                      PCR in example 7 to amplify
                                                       Length 41;
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Conservative

Mismatches

Indels

0

Gaps

GATTTCTAGGAATTCAAA 19

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RESULT 23
US-10-327-481A-19/c
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APPLICANT: King, Andrew M.O.
APPLICANT: Burman, Alison J
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NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn Ver. 2.1
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PRIOR FILING DATE: 2001-06-27
PRIOR APPLICATION NUMBER: FR 0008437
                                                                                                                                                                                SEQ ID NO 25
LENGTH: 44
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CURRENT APPLICATION NUMBER: US/10/327,481A
CURRENT FILING DATE: 2002-12-20
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Burman, Alison J.
APPLICANT: Audonnet, Jean-Christophe F.
APPLICANT: Lombard, Michel F.A.
TITLE OF INVENTION: Vaccine against Foot-and-Mouth Disease
FILE REFERENCE: 454313-3178
CURRENT APPLICATION NUMBER: US/10/327,481A
CURRENT APPLICATION NUMBER: US/10/327,481A
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APPLICANT: Lombard, Michel F.A.
TITLE OF INVENTION: Vaccine against Foot-and-Mouth Disease
            Query Match
Best Local Similarity
                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: PCT/FR01/02042
PRIOR FILING DATE: 2001-06-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: oligonuclectide JCA323 used in PCR in example 6 to amplify OTHER INFORMATION: a fragment of about 3462 bp
                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                           OTHER INFORMATION: oligonucleotide JCA329 used in PCR in example 7 to amplify OTHER INFORMATION: a fragment of about 3550 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH: 44
                                                                                                                                            ORGANISM: Artificial sequence
                                                                                                                                                                 TYPE: DNA
                                                                                                                        FEATURE:
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70.5%;
88.9%;
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Pred. No. 2.
  Score 14.8; DB 12;
Pred. No. 2.6e+03;
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                     Length 44;
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US-10-197-293-46
                                                                                                                    , OTHER INFORMATION: Oligonucleotide ZC14819 US-10-197-293-46
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Publication No. US20030171547A1
GENERAL INFORMATION:
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APPLICANT: Conklin, Darrell C.
APPLICANT: No. US200Z0058801Alak, Julia E.
TITLE OF INVENTION: Mammalian Alpha Helical Protein-1
FILE REFERENCE: 97-7101
                                                                                                                                                                                                                          SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1997-12-10 NUMBER OF SEQ ID NOS: 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 09/209,525
PRIOR FILING DATE: 1998-12-10
PRIOR APPLICATION NUMBER: 60/067,779
                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/932,679
CURRENT FILING DATE: 2001-08-16
                                                    Matches
                                                                                                                                                                                                                                       PRIOR FILING DATE: 1997-08-26
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/197,293
CURRENT FILING DATE: 2002-07-16
PRIOR APPLICATION NUMBER: US 09/686,838
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Sheppard, Paul O.
APPLICANT: Humes, Jacqueline M.
TITLE OF INVENTION: ADIPOCYTE-SPECIFIC PROTEIN HOMOLOGS
FILE REFERENCE: 97-49C1
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/140,804 PRIOR FILING DATE: 1998-08-26
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TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/056,983
                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2000-10-10
                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                         FEATURE:
                                                                                                                                                                                                         ENGTH: 58
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1 AGATTTCTAGGAATTCAAATC 21
                                                    l Similarity
17; Conserv
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                                                      Conservative
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Pred. No. 3.4e+03;
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                                                                           Score 14.6; DB 13;
Pred. No. 3.4e+03;
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RESULT 29
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CURRENT APPLICATION NUMBER: US/09/908,975
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
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                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.0 SEQ ID NO 14277
                                                                                                                           Matches
                                                                                                                                    Query Match 66.7%; Score 14, DB 13; Best Local Similarity 100.0%; Pred. No. 6.6e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14277, Application US/09908975
Publication No. US20030165843A1
GENERAL INFORMATION:
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LENGTH: 99
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                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SHOSHAN, Avi
APPLICANT: WASSERMAN, J
APPLICANT: MINTZ, Eli
APPLICANT: MINTZ, Liat
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PRIOR FILING DATE: 2001-01-13
PRIOR APPLICATION NUMBER: US 09/855,768
PRIOR FILING DATE: 2001-05-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Soybean SSRs and Methods of Genotyping FILE REFERENCE: 38-10 (52679) A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/969,373
CURRENT FILING DATE: 2001-10-02
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                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                    LENGTH:
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                                                 GATTTCTAGGAATT 39
                                                                                   GATTTCTAGGAATT 15
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84.2%;
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Sequence 115, Application US/09920033
Publication No. US20030087853A1
GENERAL INFORMATION:
APPLICANT: Rosanne M. Crooke
APPLICANT: Mark J. Graham
APPLICANT: Mark J. Graham
TITLE OF INVENTION: ANTISENSE MODULATION OF APOLIPOPROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-388-263-651/c
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                                                                                                                                                                 NUMBER OF SEQ ID NOS: 947

SOFTWARE: FRAESEQ for Windows Version 4.0

SEQ ID NO 651

LENGTH: 20

TYPE: DNA
ORGANISM: Artificial Sequence
                                                              Matches
                                                                           Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Publication No.
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SEQ ID NO 115
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Best Local Similarity
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/388,263
CURRENT FILING DATE: 2003-03-12
                                                                                                                                                                                                                                                                                                            APPLICANT: Wyatt, Jacqueline R.
APPLICANT: Borchers, Alexander
APPLICANT: Vickers, Timothy A.
TITLE OF INVENTION: IDENTIFICATION OF GENETIC TARGETS FOR
TITLE OF INVENTION: MODULATION BY OLIGONUCLEOTIDES AND
TITLE OF INVENTION: GENERATION OF OLIGONUCLEOTIDES FOR GENE MODULATION
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CURRENT APPLICATION NUMBER: US/09/920,033
CURRENT FILING DATE: 2001-08-01
                                                                                                                                                           FEATURE:
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15; Conserv
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GATTTCAAGGGATTCAA 4
                          GATTTCTAGGAATTCAA 18
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                                                             Conservative
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Henri M.
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88.2%;
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88.2%;
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Pred. No. 6.6e+03;
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Pred. No. 6.6e+03;
                                                             Mismatches
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US-10-147-196-115/c; Sequence 115, Application US/10147196; Publication No. US20030215943A1

GENERAL

INFORMATION:

RESULT 31

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FILE REFERENCE: 36688-0005
CURRENT APPLICATION NUMBER: US/09/908,975
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 60/287,724
PRIOR FILING DATE: 2001-05-02
PRIOR APPLICATION NUMBER: US 60/221,607
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
SOFTWARE: PatentIn version 3.0
SEQ ID NO 20387
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LENGTH: 20
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Publication No. US20030165843A1
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CURRENT FILING DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 124
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TITLE OF INVENTION: ANTISENSE MODULATION OF APOLIPOPROTEIN B EXPRESSION
FILE REFERENCE: ISPH-0664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rosanne M. Crooke
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APPLICANT: WASSERMAN, Alon
APPLICANT: MINTZ, Eli
APPLICANT: MINTZ, Liat
APPLICANT: WASSERMAN, Alon
APPLICANT: MINTZ, Eli
APPLICANT: MINTZ, Liat
APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0005
                                                                                            APPLICANT: SHOSHAN, AVI
APPLICANT: WASSERMAN,
APPLICANT: MINTZ, Bli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLECTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
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ORGANISM: Artificial Sequence
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Pred. No. 8.
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Pred. No. 6.6e+03;
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; ORGANISM: Mus musculus US-09-908-975-30132
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US-10-307-441-42
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 26232
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                                                                                                                                                                                                                                                                                              Sequence 42, Application US/10307441 Publication No. US20030166236A1
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Best Local
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PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
                 SEQ ID NO 42
                                                                                   APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Modified Xylanases Exhibiting Increased Thermophilicity
TITLE OF INVENTION: and Alkalophilicity
FILE REFERENCE: 027367-5006US
CURRENT APPLICATION NUMBER: US/10/307,441
CURRENT ETLING DATE: 2002-12-02
PRIOR APPLICATION NUMBER: PCT/CA01/00769
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/213,803
DBIOR STITUM NAME 2001-05-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/276,759
PRIOR FILING DATE: 2001-03-16
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CURRENT APPLICATION NUMBER: US/10/098,263B
CURRENT FILING DATE: 2003-01-08
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CURRENT FILING DATE: 2001-07-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Mittman, Michael TITLE OF INVENTION: Human Microarray
                                  PRIOR FILING DATE: 2000-05-31
NUMBER OF SEQ ID NOS: 51
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                             APPLICANT: SUNG, Wing L. APPLICANT: National Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 65
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LENGTH: 44
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Pred. No. 8.3e+03;
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Pred. No. 8.6e+03
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US-10-360-149-69/c
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 69
                                                                                                                                                                                                                                                                                                                          Sequence 69, Application US/10360149
Publication No. US20030219786A1
                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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LENGTH: 28
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Best Local
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                                               CURRENT APPLICATION NUMBER: US/10/360,149
CURRENT FILING DATE: 2003-02-06
PRIOR APPLICATION NUMBER: US/09/927,876
PRIOR FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: 60/225,035
PRIOR FILING DATE: 2000-08-11
PRIOR APPLICATION NUMBER: 60/202,724
PRIOR APPLICATION NUMBER: 60/202,724
PRIOR FILING DATE: 2000-05-08
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APPLICANT:
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Publication No.
                                                                                                                                                                                                                                   APPLICANT: El Tayar, Nabil
APPLICANT: Campbell, Rober
APPLICANT: Kelton, Christi
APPLICANT: He, Chaomei
                                PRIOR FILING DATE: 2000-05-08
NUMBER OF SEQ ID NOS: 107
                                                                                                                                                                                              APPLICANT: He, Chaomei
TITLE OF INVENTION: No. US20030219786Alel Glycoproteins and Methods
FILE REFERENCE: 20993-003
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CURRENT APPLICATION NUMBER: US/09/927,876
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: 60/225,035
PRIOR FILING DATE: 2000-08-11
PRIOR APPLICATION NUMBER: 60/202,724
PRIOR FILING DATE: 2000-05-08
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SOFTWARE: PatentIn Ver. 2.1
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TITLE OF INVENTION: No. US20040005554Alel Glycoproteins and Methods of Use Thereof
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                                                                                                                                                                                                                                                  Campbell, Robert K
Kelton, Christie A
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Kelton, Christie A
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No. US20040005554A1
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93.3%;
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Pred. No. 1.1e+04;
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Pred. No. 9
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US-10-251-117-633/c
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Publication No. US20030170891A1
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANTION: Gene Expression Using Short Interfering RNA
FILE REFERENCE: 900/042 (MBHB02-468-A)
CURRENT FILING DATE: 2003-02-24
PRIOR APPLICATION NUMBER: US 60/393,924
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 60/358,580
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SEQ ID NO 28326
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Publication No. US20030165843A1
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Best Local
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APPLICANT: FAIGLER, Simchon
TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND SPLICE
TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
FILE REFERENCE: 36688-0005
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APPLICANT: WASSERMAN,
APPLICANT: MINTZ, Eli
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PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 32337
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TYPE: DNA
ORGANISM: Mus musculus
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Similarity 93.3%;
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MINTZ, Eli
MINTZ, Liat
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Pred. No. 1
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Pred. No. 1.1e+04;
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; PRIOR APPLICATION NUMBER: US 60/296,249
; PRIOR FILING DATE: 2001-06-06
; NUMBER OF SEQ ID NOS: 1213
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 63
; LENGTH: 19
; TYPE: RNA
; ORGANISM: Artificial Sequence
; PRATURE:
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US-10-251-117-940
; Sequence 940, Application US/10251117
; Publication No. US20030170891A1
; Publication No. US20030170891A1
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Search completed: February 4, 2004, 19:45:55
Job time : 151.409 secs
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Matches 10
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Best Local Similarity 83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: RNA Interference Mediated Inhibition of Epidermal Growth Factor TITLE OF INVENTION: Gene Expression Using Short Interfering RNA FILE REFERENCE: 900/042 (MBHB02-468-A) CURRENT APPLICATION NUMBER: US 60/351,117 CURRENT FILING DATE: 2003-02-24 PRIOR APPLICATION NUMBER: US 60/393,924 PRIOR APPLICATION NUMBER: US 60/393,924 PRIOR FILING DATE: 2002-07-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 10/163,552
PRIOR FILING DATE: 2002-06-06
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR FILING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 09/916,466
PRIOR APPLICATION NUMBER: US 60/296,249
PRIOR APPLICATION NUMBER: US 60/296,249
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NUMBER OF SEQ ID NOS: 1213
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Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

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                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP)
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                                                                                                                                                                                                                                                                                                                     ВJ044866
                                                                                                                                                                                                                                                                                                                                     BZ352520
                                                                                                                                                                                                                                                                                                                                                             TA383F04Q
                                                                                                                                                                                  ALIGNMENTS
                                                                       Catarrhini; Hominidae;
                                                                                Craniata; Vertebrata; Euteleostomi;
                                                                                                                                    Homo sapiens
                                                                                                                                            mRNA
                                                                                                                                             linear
                                                                                                                                     cDNA clone
                                                                                                                                                                                                                                        AL632588 AL632588
BZ662433 SALK 0259
BH909538 SALK 0543
BZ353011 SALK 1196
BZ355066 SALK 1263
                                                                                                                                                                                                                                                                                                                                                                                           W09386 ma63e02.rl
AF087410 AF087410
CC044689 3591_1_16
CC044746 3591_1_16
CC457081 SALK_1061
                                                                                                                                                                                                                                                                                                                                                                                                                                          BE151753 QV0-HT030
BE762761 QV3-NT002
BM447217 DSA007D11
BM738230 K-EST0002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AW582100 MR3-ST019
AW781526 sl81c04.y
AA513622 nh29e11.s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA015156 mi63a07.r
AW101871 sd71g06.y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AW459607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BZ663770 SALK 0273
BE248200 NF002G04D
BX012889 Single re
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AZ592186 1M0402H24
BZ663770 SALK_0273
                                                                                                                                                                                                                  BZ762011 SALK 0842
AL937418 Arabidops
AI307980 ta89h09.x
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BQ287277 ii65e03.x
                                                                                                                                                                                                           AU007620 AU007620
                                                                                                                                                                                                                                                                                                                                                                                  BX002146 Arabidops
                                                                                                                                                                                                                                                                                CC459429 SALK_1297
AG219794 Lotus jap
                                                                                                                                                                                                                                                                                                                                                              AL482493 T. brucei
AL497990 T. brucei
                                                                                                                                                                                                                                                                                                                                                                                                                                   BW069148 BW069148
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MR3-ST019
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                                                                                                                                                                                                                                                                                                                                                     1M0553B07
                                                                                                                                                                                                                                                                                                2M0197K12
                                                                                                                                                                                                                                                                                                        1M0042F02
                                                                                                                                                                                                                                                                                                                       BJ044866
                                                                                                                                                                                                                                                                                                                               SALK
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97 bp mRNA linear EST 04 NISC 1909d11.y1 NCI CGAP Pr32 Rattus norvegicus cDNA clone IMAGE:5622956 5', mRNA sequence.
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Seq primer: -40UP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov

cDNA Library Preparation:

cDNA Library Arrayed by: The I.M.A.G.E. Consortium/LLNL

DNA Sequencing by: National Institutes of Health Intramural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus
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                                                                                                                                                                                                                                                                                                                                                         Plate: LLAM12449 row: G column: Seq primer: M13RP1 reverse primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequencing Center (NISC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus (Norway rat)
                                                                                                                                                                                                                                                                                                                                                                                                           info@image.llnl.gov
Plate: LLAM12449 ro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clone distribution: NCI-CGAP clone distribution information can ound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAATTTCTAGGAATTAAAAGC
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Location/Qualifiers
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/lab_host="DH10B"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"
/tissue_type="dorsal prostate"
/dev_stage="adult, 14 month"
                                                                                                                           /clone="IMAGE:5622956"
                                                                                                                                                                                                             Tom
                                                                                                                                                                                                                                              organism="Rattus norvegicus"
                                                                                                                                                                                                                                                                                                                                 location/Qualifiers
                                                                                      sex="male"
                                                                                                                                                                    db_xref="taxon:10116"
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85.7%;
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Pred. No. 5.1e+03;
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Location/Qualifiers
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Plate: 0023 row: P column:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse whole genome scaffolding with paired end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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AZ782789.1 GI:12916863
GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Utah
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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. USA
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    adaptored DNA was purified and size-selected for a 9.5 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivate
                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F/clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                    Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab host="DH10B (Tl phage-resistant)"
/clone_lib="NCI_CGAP_Pr32"
/note="Torgan: prostate; Vector: pCMV-spORT6; Site_1: NotI;
/site_2: SalI; Cloned unidirectionally. Primer: Olīgo dT.
Average insert size 2 kb. Constructed by Invitrogen. Note:
this is a NCI_CGAP_Library."
a 15 c 12 g 15 t
                                                                                                                                                                                                                                                                                                                                                                       mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
                                                                                                                                                                                                                                                                                                               sex="Male"
                                                                                                                                                                                                                                                                                                                               clone="UUGC2M0023P07"
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85.0%;
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Pred. No. 1.4e+04;
0; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Std Error: 0.00
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  DNA was prepared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA
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from a derivative
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VERSION
KEYWORDS
SOURCE
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COMMENT
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CB405258
                                                                                       BASE COUNT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  designed on the predicted protein encoding ORF. C. elegan cloning project: Contact david_hill@dfci.harvard.edu or marc_vidal@dfci.harvard.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marc Vidal Laboratory
Dana Farber Cancer Institute
1 Jimmy Fund Way Smith 858, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ,C.M., Li,S., Jacotot,L., Bertin,N., Janky,R., Moore,T., Hudson,J.R., Hartley,J.L., Brasch,M.A., Vandenhaute,J., Boulton,S., Endress,G.A., Jenna,S., Chevet,E., Papasotiropoulos,V., Tolias,P.P., Ptacek,J., Snyder,M., Huang,R., Chance,M.R., Lee,H., Doucette-Stamm,L., Hill,D.E. and Vidal,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence tag of Gateway entry clones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 617 632 5180 Fax: 617 632 5739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nat. Genet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reboul, J., Vaglio, P., Rual, J.F., Lamesch, P., Martinez, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OSTR040A2_1 AD-wrmcDNA Caenorhabditis elegans cDNA, mRNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                 POLYA=No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: Marc_Vidal@dfci.harvard.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Vidal M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Caenorhabditis elegans
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                                                                                       24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                         /note="The AD wriceNA library was generated with poly(A)+
NNA isolated from both hermaphrodite and male N2 worms of
all larval stages, embryos, adults and dauers and the
subsequent generation of cDNAs by poly(A) priming. The
cDNAs were cloned into pPC86"
a 19 c 19 g 25 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

18 c 14 g 28 t
                                                                                                                                                                                                                                /sex="Hermaphrodite and male"
/tissue_type="whole animal"
/dev_stage="mixed stage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                     æ
                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                Clone_lib="AD-wrmcDNA"
                                                                                                                                                                                                                                                                                                                                                                organism="Caenorhabditis elegans"
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                                                                                                                                                                                                                                                                                                 xref="taxon:6239"
                                                                                                                                                                                                                                                                                                                                       _type="mRNA"
72.4%;
85.0%;
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0; Mismatches
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Pred. No. 1.4e+04;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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Query Match Best Local Similarity

Score Pred.

No. 1.5e+04;

Length 87;

rosids

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RESULT 5
AL940051/c
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                                                                                                                                                                              ORIGIN
                                                                                                                                                                                               BASE COUNT
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TITLE
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                                                                                                                     Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
                             67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone F1809. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line availability can be found at:

http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSS
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AL940051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana T-DNA flanking sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rosso, M., Strizhov, N., Li, Y. and Weisshaar, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rosso, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and Weisshaar, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for flanking sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
                                                              1 AGATTTCTAGGAATTCAAAT 20
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                                                                                                                       Similarity
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                             AGTTTTCGAGGAATTCAAGT 48
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                                                                                                                                                                                                                                 /clone lib="Arabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
flanking the insertion. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
processed for submission. T-DNA derived sequences were
                                                                                                                                                                                                 removed"
a 19 c
                                                                                                                                                                                                                                                                                                                                                                                                                       /mol_type="genomic DNA"
/strain="Columbia 0"
/db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Arabidopsis thaliana"
                                                                                                                                                                                                                                                                                                                                                                                                           /clone="GK-243H10-014396"
                                                                                                                       72.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        v,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. thaliana T-DNA mutagenised population (GABI-Kat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rosso, M., Viehoever, P.,
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                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tag based reverse genetics
                                                                                                                       Score 15.2; DB 29;
Pred. No. 1.5e+04;
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GK-243H10-014396,
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T095D05 Populus apical shoot cDNA library Populus tremula x Populus
tremuloides cDNA 5 prime, mRNA sequence
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Populus tremula x Populus tremuloides
Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
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Department of Plant Physiology
University of Umea, 901 87 Umea,
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Fax: +46 90 786 6676
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
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                                                                                                                                                                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids I; Malpighiales; Salicaceae; Populus.
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The poplar tree transcriptome: Analysis of expressed sequence tags
                                                                                                                                                                                                                                                                                                                                                                           Populus tremula
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                                                                          Tel: +46 90 786 5279
Fax: +46 90 786 6676
Email: rupali.bhalerao@plantphys.umu.se
                                                                                                                                 Department of Plant Physiology University of Umea, 901 87 Umea,
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/clone lib="Populus apical shoot cDNA library"
13 c 16 g 32 t
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/db_xref="taxon:47664"
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/mol_type="mRNA"
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                     organism="Populus tremula"
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                                                                                                                                                                                                                                                                                                                                                                                                       CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consorthum/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Louis M. Staudt, M.D., ph.D.,
Ph.D., Gerald Marti, M.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                     Insert Length:
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                                                                                                                                                                                                                                                                                                    /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                      tissue_type="germinal center B cell"
                                                                                                                                                                                                                                                                                       clone="IMAGE:1309206"
                                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
|
| mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                            ocation/Qualifiers
                                                                                                                                                                                                                                                              lab host="DH10B"
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/clone_lib="Populus petioles cDNA library"
16 c 14 g 32 t
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 69.5%;
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Score 14.6; DB 9;
Pred. No. 2.7e+04;
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Pred. No. 1.5e
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        Length 73;
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VERSION KEYWORDS

AW695026.2

GI:11957373

ACCESSION

NF082E06ST 5', mRNA sequence AW695026

SOURCE

EST. Medicago truncatula

(barrel medic)

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NF082E06ST1F1050 Developing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10010 N. Torrey Pines Road,
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alonso, J.M., Leisse, T.J., Harajas, P., Chen, H., C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutatic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The Salk Institute Genomic Analysis Laboratory
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arabidopsis Genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Joseph R. Ecker
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93 bp DNA linear GSS 07-040
SALK 070370.34.05.n Arabidopsis thaliana TDNA insertion lines
Arabīdopsis thaliana genomic clone SALK_070370.34.05.n, genom
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BZ593365
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                                                                                                                                                                                                                                                        /Clone libe"Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:3702"
/clone="SALK_070370.34.05.n"
                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="genomic
/strain="Columbia (
                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Arabidopsis thaliana"
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                                                                                                                                                                                      69.5%;
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Pred. No. 2.8e+04;
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truncatula cDNA clone
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Prednis, L.,
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CA513671/c
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids
spermatophyta; Fabales; Fabaceae; Papilionoideae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        He,X.-Z., Shadle,G., Scott,A.D., Harris,A.R., Gonzales,R.A., Be, C.J., Flores,H.R., Inman,J.T., Weller,J.W., May,G.D. and Dixon,R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plant Biology Division
The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway, Ardmore, OK 73402, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished
On Apr 14, 2000 th
Contact: Dixon RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30
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Fax: 580 221 7380
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                                                                                                                                                                                                                                                                                                                                 10-3 1H2 Porcine testis phage
10-3 1H2, mRNA sequence.
CA513671
CA513671.1 GI:25014228
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                                                                                                                                                                                                                                                                                                                    EST.
Email: burkin@uiuc.edu
Insert Length: 93 Std Error:
Plate: 10-3_1 row: H column:
High quality sequence stop: 93.
Location/Qualifiers
1: .93
                                                                                             University of Illinois at Urbana-Champaign
1207 W Gregory Dr, Urbana, IL 61801, USA
Tel: 217 244 5144
Fax: 217 333 8286
                                                                                                                                                                                                1 (bases 1 to 93)
Burkin, H.R., Vieira, A.P. and Miller, D.J.
Direct submission (Burkin, Vieira, Miller)
                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                   Sus scrofa
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                                                                                                                                                                   Contact: Burkin HR
                                                                                                                                                                                Unpublished
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te: 082 row: E column: 06
primer: TCACACAGGAAACAGCTATGAC.
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/clone_lib="Developing stem"
/note="Vector: Lambda Zap; Contains
internodal stem segments"
16 c 5 g 11 t
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/db_xref="taxon:3880"
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nage library Sus scrofa cDNA clone
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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 97)
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GSS.
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University of Utah
Rm. 308, Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 97.
                                                                                                                                                                                                                                                                                                                                                                   Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0225 row: B column: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             84112,
                                                                                                                                                                                                                                                                                                                                                      Class: plasmid ends
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/db_xref="taxon:9823"
/clone="10-3_1H2"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone lib="Porcine testis phage
/note="Organ: testis; Vector: T7
Eco RI; Site_2: HindIII"
24 c 17 g 29 t
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                                                                                                  /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jacksor
laboratory Mouse DNA Resource
                was blunt end-repaired with T4 DNA polymerase and T4 was blunt lend-repaired with T4 DNA polymerase and T4 polymerase and T4 polymerase.
                                                                                                                                                                                                                                              /mol_type="genomic_DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                               organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                    ocation/Qualifiers
                                                                                  (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                                                                               'clone="UUGC1M0225B12"
                                                                                                                                                                                                                              'db_xref="taxon:10090"
ligated to the blunt ends in high
                                                                                                                                                                                             sex="Male"
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93.8%;
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Pred. No. 3.4e+0
0; Mismatches
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  ery Match
t Local Similarity
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Tel: 858
                                                                                                                                                                                                                                                                                                                                                                                                                   This is single pass sequence recovered from the left border of TDNA. This sequence lies within 300 bases of the 5' end of At5g16310 and 300 bases of the 5' end of At5g16320.
                                                                                                                                                                                                                                                                                                                                                                                                             Class: TDNA tagged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 48)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H.,
Zimmerman,J. and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ecker@salk.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          48 bp DNA linear GSS 13-JUN-2002 SALK 074019.36.90.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK 074019.36.90.x, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis Genome
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Arabidopsis thaliana
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                                                              /clone lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 4 c 12 g 20 t
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67.6%;
84.2%;
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Pred. No. 3.4e+04;
0; Mismatches 1;
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Score 14.2; DB 28; Pred. No. 3.8e+04;

Length 48,

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EST.
Glycine max (soybean)
Glycine max
                                                                                                                                                                                                                         20 GAATTCGAGGAATTCAAGT
                                                                                                      83 bp
sc36h04.yl Gm-c1014 Glycine max
Gm-c1014-1232 5', mRNA sequence.
AI966296
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
                                                                                                                                                AI966296
                                                                                                                                                                                                                                                  2 GATTTCTAGGAATTCAAAT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium/LLNL
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammaru, (bases 1 to 60)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nag58g08.x1 NCI_CGAP_Co26 Homo sapiens cDNA clone IMAGE:4225743
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                                                                                                                                                                                                                                                                                                                                                  colonic mucosa, cDNA made by oligo-dT priming.
Directionally cloned into UDG sites. Size-selected on agarose gel, average insert size 300 bp. Primary library.
CDNA Library Preparation: David B. Krizman, Ph.D.
Reference: Krizman et al. (1996) Cancer Research
56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="IMAGE:4225743"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tissue_type="normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
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84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                       RESULT 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shoemaker,R., Keim,P., Vodkin,L., Erpelding,J., Coryell,V., Khanna,A., Bolla,B., Marra,M., Hiller,L., Kucaba,T., Martin,J., Beck,C., Wylie,T., Underwood,K., Steptoe,M., Theising,B., Allen,M., Bowers,Y., Person,B., Swaller,T., Gibbons,M., Pape,D., Harvey,N., Schurk,Y., Ritter,E., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae; Glycine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Shoemaker R/Public Soybean EST Project public Soybean EST Project Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                           12 AAATTTCAAGGAATTCGAA 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              call: (800)-533-4363 or contact via email: ccu@resgen.com Seq primer: -40RP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This clone is available through: ResGen, Invitrogen Corp. South Memorial Parkway Huntsville, AL 35801 for further in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
Tel: 314 286 1800
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                                                                                                                                                                                                                      47 bp mRNA linear BST 11-DEC-:
BJ077154 NIBB Mochii normalized Xenopus tailbud library Xenopus
laevis cDNA clone XL060a15 3', mRNA sequence.
BJ077154
BJ077154
BJ077154.1 GI:17522070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                   EST.
                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea;
Xenopodinae; Xenopus.
1 (bases 1 to 47)
                                                                                                                                                                                        Xenopus laevis (African clawed frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                   Xenopus laevis
                                               Kitayama,A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 83)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGATTTCTAGGAATTCAAA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone lib="Gm-c1014"
/note="Wector: pT/TJ3Pac (pT/TJ, pharmacia); Site 1: EcoRI;
/note="Wector: pT/TJ3Pac (pT/TJ, was constructed from
Site 2: HindIII; This cDNA library was constructed from
mRNA isolated from leaves of 2-3 week old greenhouse grown
plants. Complementary DNA was synthesized from mRNA using
plants. Complementary DNA was synthesized from mRNA using
plants to the blunt-ended cDNA fragments followed by
ligated to the blunt-ended cDNA fragments were
digestion with EcoRI and HindIII. The cDNA fragments were
directionally cloned into the EcoRI-HindIII restriction
directionally cloned into the bigated cDNA fragments
site of the pT/T3-Pac vector. The ligated cDNA fragments
were transformed into DH10B host cells (Gibco BRI). This
were transformed into DH10B host cells (Gibco BRI). This
library was constructed by Dr. Randy Shoemaker and Dr.
John Erpelding."

32 a 9 c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Glycine max"
/mol_type="mRNA"
/db_xref="taxon:3847"
/clone="GENOME SYSTEMS CLONE ID: Gm-c1014-1232"
/tissue_type="Leaves, 2-3 week old seedlings, greenhouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       location/Qualifiers
genes in X. laevis embryo
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                                             Terasaka, C., Mochii, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.6%;
84.2%;
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Pred. No. 4.1e+04;
0; Mismatches 3;
                                                    Ueno, N., Shin-i, T. and Kohara
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                                                                                                                                   Buteleostomi;
; Pipidae;
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COMMENT
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AA422541
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Center For Genetic Resource Information
National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Tadasu Shin-i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 49)

1 (bases 1 to 49)

Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and Waterston P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vf14c03.s1 Knowles Solter mouse unfertilized clone IMAGE:835684 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                        Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA422541.1
                                                                                                                                                                                                                                                                                                                                                   Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                  The WashU-HHMI Mouse EST Project
                                                                                                                                                                                                                      Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                      Fax: 314 200 1000 Email: mouseest@watson.wustl.edu
Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LINL; contact the
                                                                                                                                                                                                                                                                                                                    Fax: 314 286 1810
                                                                                                                                                                                                                                               MGI:495900
                                                                                                                                                                                                                                                            IMAGE Consortium (info@image.llnl.gov) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGTTCTAGGAATCCAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTTCTAGGAATTCAAA 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Xenopus laevis"
/mol_type="mRNA"
/db_xref="taxon:8355"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="whole embryo"
/dev_stage="stage 25"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="XL060a15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone
/clone_lib="Knowles Solter mouse unfertilized egg"/note="Organ: unfertilized egg; Vector: pBluescribe (modified); Site_1: MluI; Site_2: SalI; Cloned
                                                       /tissue_type="unfertilized
/lab_host="DH10B"
                                                                                                                                                                   organism="Mus musculus"
                                                                                                                                                                                                        ocation/Qualifiers
                                                                                            /clone="IMAGE:835684"
                                                                                                            db_xref="taxon:10
                                                                                                                                   strain="C57BL/6J x DBA/2J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.7%;
88.2%;
                                                                                                                                                  type="mRNA"
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Pred. No. 5.
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AZ810918/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0076 row: K column: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished
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Mus musculus
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             quality sequence stop: 69
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                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                       clone
                                                                                                                                                                                                                                                                                                              sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     eggs. Primer: Sall(dT):
5'-CGGTCGACCGTCGACCGTTTTTTTTTTTTT-3'. cDNAs were
cloned into the Mul/Sall sites of a modified pBluescribe
vector using commercial linkers (NEB). Average insert
size: 1.0 kb."
                                                                                                                                                                                                                               lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"clome lib="Mouse 10kb plasmid UUGCIM library" clome lib="Wouse 10kb plasmid UUGCIM library" note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                              clone="UUGC2M0076K14"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             unidirectionally from mRNA prepared from 5000 unfertilized
                                                                                                                                                                                                                                                                                                                                                      _xref="taxon:10090"
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C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.8; DB 9;
Pred. No. 5.8e+04;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Murinae; Mus.
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           High quality sequence stop: 69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0223 row: C column: 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 TTTCTAGGAATTCAAAT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                308,
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                                                                                          /lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse_10kb_plasmid_UUGC2M_library"
/note="Vector: PMD42nv; Purified_genomic_DNA_from_M.
musculus_C57BL/6J_Remale) was obtained_from_the_Jackson_resistant, F-"
Taborestor; Manager Name Taborestory, Name
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                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Mus musculus"
/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex≈"Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'db_xref="taxon:10090"
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'strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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88.2%;
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No. 6.1e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA
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E., Pedersen,T., Reilly
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BG153467/c
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AUTHORS
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1 (bases 1 to 76)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BG153467.1 GI:12665497
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24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   info@image.llnl.gov
Seq primer: -40UP fo
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                                                                                              Similarity
GAATTCGAGGAATTCAA
                                        GATTTCTAGGAATTCAA 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

19 c 11 g 21 t
                                                                                                                                                                                                               /clone lib="NCI_CGAP_CO29"
/note="Organ: colon; Vector: pAMP1; mENA made from colonic /note="Organ: colon; Vector: priming. Directionally adenoma, CDNA made by oligo-dT priming. Directionally cloned into UDG sites. Size-selected on agarose gel, cloned into UDG sites. Size-selected on agarose gel, average insert size 300 bp. Primary library. cDNA Library preparation: David B. Krizman, Ph.D. Reference: Krizman preparation: David B. Krizman, Ph.D. Reference: Krizman
                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                              tissue type="tubulovillous adenoma"
                                                                                                                                                                                                                                                                                                                                                                    clone="IMAGE:4204320"
                                                                                                                                                                                           al. (1996) Cancer Research 56:5380-5383
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88.2%;
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                                                                                                Score 13.8; DB 10; Pred. No. 6.2e+04;
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BG151856/c
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                                   JOURNAL
                                                                                      AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium/LLNL
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BG151684
BG151684.1 GI:12663714
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
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                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 88)
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                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                             Homo sapiens (human)
              Contact: Robert Strausberg, Ph.D.
                                   Unpublished
                                                     Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         colonic mucosa, cDNA made by oligo-dT priming.
Directionally cloned into UDG sites. Size-selected on agarose gel, average insert size 300 bp. Primary library.
CDNA Library Preparation: David B. Krizman, Ph.D.
Reference: Krizman et al. (1996) Cancer Research
56:5380-5383."

14 g 24 *
cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="normal colonic mucosa"
/lab_host="DH10B"
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88.2%;
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Pred. No. 6.2e+04;
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                                                                                                                                                                                                                                                                                          dq 88
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                                                                                                                                                                                                                                                                         p mRNA linear EST 05-FEB-2
sapiens cDNA clone IMAGE:4226360
                                                                           Genome Anatomy Project
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BX533932
                FEATURES
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availability can be found at:
http://www.mpiz-koeln.mpg.de/GABI-Kat/.
Location/Qualifiers
                                                            Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone F21J9. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                35 GAATTCGAGGAATTCAA 19
                                                                                                                                                                                                                                     3 (bases 1 to 88)
Strizhov, N., Li, Y., Rosso, M. and Weisshaar, B.
Direct Submission
                                                                                                                                                                                                      Submitted (02-JUN-2003) Weisshaar B., Max-Planck-Institut fuer
                                                                                                                                                                                                                                                                                                                                              Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat)
                                                                                                                                                                                                                                                                                                         Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                 for flanking sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and Weisshaar, B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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Seq primer: -40UP f1
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/note="Organ: colon; Vector: pAMP1; mRNA made from normal
/note="Organ: colon; Vector: pAMP1; mRNA made from normal
colonic mucosa, cDNA made by oligo-dT priming.
Directionally cloned into UDG sites. Size-selected on
agarose gel, average insert size 300 bp. Primary library.
cDNA Library Preparation: David B. Krizman, Ph.D.
Reference: Krizman et al. (1996) Cancer Research
56:5380-5383."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="normal colonic mucosa"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="IMAGE:4226360"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:9606"
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88.2%;
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                                                                                                                                                                                                                                                                                                                          tag based reverse genetics
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Pred. No. 6.3e+04;
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Best Local :
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                                                                                                                                                                                                                                                                                   Plate: mgmy001
Seq primer: T3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Ebbole DJ
Department of Plant Pathology & Microbiology
Texas A&M University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peterson Bldg, MS2132,
Tel: 979 845 4831
Fax: 979 845 6483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mgmy001xJ18f.b Magnaporthe grisea MY Uni-Zap XR Library Magnaporthe grisea cDNA clone mgmy001xJ18 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                               BACKWARD: T7
                                                                                                                                                                                                                                                                                                                                                 FORWARD: T3
                                                                                                                                                                                                                                                                                                                                                                                                            Chromatogram file of this sequence is available,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 91)
Ebbole, D.J., Yuan, J., Thomas, T.L., Bobrowicz, P., Lu, G., Bhatterai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Magnaporthe grisea
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                                                                                                                                                                                                                                                                                                                                                                         PCR PRimers
                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: d-ebbole@tamu.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Expressed sequence tags from the rice blast fungus, Magnaporthe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Magnaporthe grisea (anamorph: Pyricularia grisea)
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/cell_type="mycelium"
/clone_lib="Magnaporthe grisea MY Uni-Zap XR Library"
/clone_lib="Magnaporthe grisea MY Uni-Zap XR Library"
/note="Vector: pBluescriptSK+; Site_1: EcoRI; Site_2:
; Unidirectional cloning. EcoRI side has T3 primer and
predominantly 5' reads. T7 primer on XhoI side of inse
                                                                                          /clone="mgmy001xJ18"
/sex="Mat1-2 hermaphrodite"
/cell_type="mycelium"
                                                                                                                                                        /db_xref="taxon:148305"
                                                                                                                                                                              /mol_type="mRNA"
/strain="70-15"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone lib="Arabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
                                                                                                                                                                                                                   organism="Magnaporthe grisea"
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/mol_type="genomic DNA"
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Pred. No. 6.3e+04;
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RESULT 25
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BH758017
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AF087391 91 bp DNA linear GSS 18-APR-:
AF087391 Chlamydia trachomatis L2 Chlamydia trachomatis genomic
clone 830 similar to RAD1 protein, genomic survey sequence.
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AF087391.1
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Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.
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Chlamydia trachomatis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        635 Barnhill Dı
Class: shotgun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Wang L
Department of Microbiology & Immunology
Indiana University School of Medicine
635 Barnhill Drive; MS 255, Indianapoli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wang,L., Steenburg,S.D., Zheng,Y. and Larsen,S.H. Gene identification of Chlamydia trachomatis by r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                       91 bp DNA linear GSS SALK 013098.27.85.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK 013098.27.85.x,
         Arabidopsis thaliana (thale cress)
Eukaryota; Viridiplantae;
                                                            BH758017.1 GI:19042773
                                                                                        survey sequence.
                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Minimal medium mycelium library. Sequences were processed by one of two methods. Where a full-length alignment to the M. grisea genome sequence was available, the EST sequence was trimed according to the alignment, otherwise sequence quality was assessed using phredPhrap version 991019 and trimmed according to phd files (0.05) and for
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/note="isolate=434B"
18 c 11 g 29 t
                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="genomic_DNA"
/strain="L2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                           'clone="830"
                                                                                                                                                                                                                                                                                                                                                                                                                          db_xref="taxon:813"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Chlamydia trachomatis"
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88.2%;
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88.2%;
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Pred. No. 6
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Pred. No. 6.3e+04;
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Streptophyta; Embryophyta; Tracheophyta;
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Joseph R. Ecker
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ecker@salk.edu
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                                                                                                                                                                                                                               Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 48)
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Drosophila melanogaster
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GSS.
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Sequence recovery method was 
Sequence orientation is forwa
                                                                                                                                   Unpublished
Contact: Gerald Rubin
                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                       lines Drosophila melanogaster genomic Sequence recovered end of P element, genomic survey sequence.
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                                                        LSA Building, Berkeley, CA 94720-3200, Fax: 5106439947
                                                                                        Berkeley Drosophila Genome Project
University of California, Berkeley
                                    Email: gerry@fruitfly.berkeley.edu
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Zimmerman,J. and Ecker,J.R.
Sequence-Indexed Library of Insertion Mutations in
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/clone=Iib="Arabidopsis thaliana TDNA insertion lines"
/clone=FCR was performed on Arabidopsis thaliana lines
mote=FCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 12 c 18 g 27 t
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/strain="Columbia 0"
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                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Jeffrey E. Green, M.D.
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           602105183F1 NCI_CGAP_Kidl4 Mus musculus cDNA clone IMAGE:4223229 5', mRNA sequence.
BF789276
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National Institutes of Health, Mammalian Gene Collection (MGC)
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The P element insertion position is base 1 in the 48 bases. This
insertion position refers to the first base of the 8 base target
recognition sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
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           /lab_host="DH10B (TI phage-resistant)"
/clone lib="NCI CGAP Kid14"
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/note="Tota="NCI CGAP Kid14"
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library. |"
a 9 g 15 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://www.fruitfly.org/about/methods/inverse.pcr.html."
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                                                                                                                                                              /db_xref="taxon:10090"
/clone="IMAGE:4223229"
                                                                                                                                                                                                         /mol_type="mRNA"
/strain="FVB/N"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: mEST@mail.nih.gov

The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to identify it as a clone from the retina tissue cDNA Library Preparation: M.B. Soares Lab Clone firm RESEARCH GENETICS. It should be noted that Bento Soares is generating a small number of additional specialized non-redundant arrays of BMAP cDNAs whose availability will be considered under appropriate and limited collaborative arrangements The following repetitive elements were found in this cDNA sequence: 15-105, >B2#SINE/B2
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Fax: 301 443 9890
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Bonaldo, M.F., Lennon, G. and Soares, M.B.
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                                                                                      /clone="UI-M-CGOp-bnp-a-11-0-UI"
//lab host="DH10B (Life Technologies)"
/clone="lib="NIH_BMAP_Ret4_S2"
/clone_lib="NIH_BMAP_Ret4_S2"
/clone_lib="Vector: pT773D-Pac (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; The NIH_BMAP_Ret4_S2 library is a subtracted library, ultimately derived from mouse retina tissue libraries at various stages of development. For a detailed description of the library from which this clone was derived, please visit our web site at brainest.eng.uiowa.edu. The tissue for this library was contributed by Dr. Xin-Yuan Fu, Yale University School of Medicine
TAG_SEQ=GTCAGCGCGCCC"
20 c 13 g
                                             TAG_LIB=NIH_BMAP_Ret4_S2
TAG_TISSUE=adult-retina
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/strain="C57BL/6J"
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Pred. No. 7.1e+04;
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Identification of gene expression profile of dorsal root ganglion in the rat peripheral axotomy model of neuropathic pain proc. Natl. Acad. Sci. U.S.A. 99 (12), 8360-8366 (2002)
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Institute of Neuroscience
320 Yue Yang Road, Shanghai 200031, P.R.China
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This clone is also available at Chinese National Human Genome Center at Shanghai, 351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong New Area, P.R.China. Please contact with Zhang Xu (xu.zhang@ion.ac.cn) or Han Zeguang (hanzg@chgc.sh.cn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Zhang Xu
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/dev_stage="adult"
/clone_lib="Rat_DRG_Library"
/clone_lib="Rat_DRG_Library"
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                                                                                                                                                                                                                                                                                                     /strain="Sprague-Dawley"
/db_xref="taxon:10116"
/clone="DRACQH10"
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/mol_type="mRNA"
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Pred. No. 7.5e+04;
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AGB DP MRNA linear EST 29-APR-1998 og91c09.sl NCI CGAP Kid5 Homo sapiens cDNA clone IMAGE:1455664 3' similar to gb:M12529 APOLIPOPROTEIN E PRECURSOR (HUMAN);, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 AGATTTCTAGGAATTCAAAT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Umea Plant Science Center
Department of Plant Physiology
University of Umea, 901 87 Umea,
                                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 86) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP), National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: rupali.bhalerao@plantphys.umu.se.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unneberg, P., Bhalerao, R.R., Jansson, S. and Sterky, F. The poplar tree transcriptome: Analysis of expressed
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1 (bases 1 to 81)
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BU890627
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P039E04 Populus petioles
                                                                 cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the L.N.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                  Unpublished Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                   AA863156.1
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             Insert Length: 486 Store Seq primer: -40ml3 fwd.
                  Tissue Procurement: Christopher Moskaluk, M.D., Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                       Tumor Gene Index
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/clone_lib="Populus petioles
24 c 20 g 21 t
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/mol_type="mRNA"
Location/Qualifiers
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Pred. No. 7.7e+04;
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                                                                                                            Sequencing Center
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eudicots; rosids
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RESULT 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87 bp DNA linear GSS 13-MAR-2003
SALK 110771.31.45.x Arabidopsis thaliana TDNA insertion lines
Arabidopsis thaliana genomic clone SALK_110771.31.45.x, genomic
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                                                                                                                                                                                                                                                                                                                                                                         Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gr,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Zimmerman,J. and Ecker,J.R.

A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                      Tel: 858 453 4100 x1752 Fax: 858 558 6379
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Joseph R. Ecker
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                                                                                                                                                                                                                                                               TDNA tagged
                                                                                                                                                                                                                                                                                                                      ecker@salk.edu
/clone="SALK_110771.31.45.x"
/clone lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
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                                                                                                                                          /db_xref="taxon:3702"
                                                                                                                                                                strain="Columbia 0"
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                                                                                                                                                                                                organism="Arabidopsis thaliana"
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Pred. No. 7
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7.8e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone F9H3. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A pipeline for automated (flanking sequence tags)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (21-OCT-2002) Weisshaar B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat) for flanking sequence tag based reverse genetics
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                                                                                                                                                                                                                                                                                                                                                                                                     http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                             availability can be found at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 (bases 1 to 90)
Rosso, M., Li, Y., Strizhov, N. and Weisshaar, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
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                 removed"
a 11 c
                                                     were directly sequenced to determine the genomic sequence flanking the insertion. Sequences displaying significant similarity to the A. thalaina nuclear genome sequence were processed for submission. T-DNA derived sequences were
                                                                                                                                         /note="PCR was performed on DNA from Arabidopsis thaliana plants (T1) which were transformed with the T-DNA from vector pAC161. The lines contain one or more T-DNA insertions. The DNA fragment(s) resulting from the PCR
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/clone_lib="Arabidopsis thaliana_T-DNA insertion lines"
                                                                                                                                                                                                                                                                          db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                            /mol_type="genomic D
/strain="Columbia 0"
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/mol_type="genomic DNA"
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Pred. No. 7.8e+04;
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Query Match
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Matches 16; Conserv

Conservative

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Score 13.6; DB 29 Pred. No. 7.8e+04; 0; Mismatches 4

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RESULT 35
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                                                                                                                                                            Caenorhabditis elegans
Caenorhabditis elegans
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida;
Rhabditidae; Peloderinae; Caenorhabditis.
1 (bases 1 to 92)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Umea Plant Science Center
Department of Plant Physiology
University of Umea, 901 87 Umea,
Tel: +46 90 786 5279
Fax: +46 90 786 6676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid; eurosids I; Malpighiales; Salicaceae; Populus.

1 (bases 1 to 91)
Unneberg, P., Bhalerao, R.R., Jansson, S. and Sterky, F.
The popular tree transcriptome: Analysis of expressed sequence tags
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P034A07 Populus petioles
                            Genome Biology Lab.
National Institute of Genetics
Yata 1111, Mishima, Shizuoka 411, Japan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
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                                                                                Contact: Yuji Kohara
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81-559-81-6855
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Location/Qualifiers
                                                                                                                an expression map
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:113636"
/tissue_type="petioles"
/clone_lib="Populus_petioles cDNA_library"
/25 c 22 g 26 t
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Spermatophyta; Magnoliophyta; Liliopsida;
clade; Panicoideae; Andropogoneae; Zea.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Class: transposon-tagged.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: walbot@stanford.edu
Plate: 1006077 row: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 650 723 2227 Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Department of Biological Sciences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
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30
                                                  /clone_lib="1006 - RescueMu Grid G"
/clone="Organ: leaf; Vector: RescueMu (engineered from
/note="Organ: leaf; Vector: RescueMu (engineered from
/note="Organ: leaf; Vector: RescueMu (engineered from
pBlueScript backbone); Site 1: BamHI; Site 2: BgIII;
RescueMu is a 4.9 kb, modified maize Mu transposon
designed to allow plasmid rescue from total genomic DNA.
Mu elements insert preferentially into transcription
units. For more information on RescueMu, go to the web
site 'www.ramdb.iastate.edu' and follow the links for
'RescueMu.' Grid G was grown at Stanford in 2000. DNA was
extracted from leaf punches, double digested using BamHI
and BgIII, and ligated to form circular plasmids. DHIOB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="hermaphrodite"
/dev_stage="embryo"
/clone_lib="Yuji Kohara unpublished cDNA:Strain N2
/ernaphrodite embryo"
hermaphrodite embryo"
1 20 c 19 g 31 t
  ampicillin."
a 8 c
                                  and BglII, and ligated to form circular plass cells were transformed and then screened on
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/strain="N2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:6239"
/clone="yk107d2"
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                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                          dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                     organism="Zea mays"
                                                                                                                                                                                                                                                                        lab_host="DH10B"
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    RescueMu

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Pred. No. 7.8e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA Library construction by: Joe Barnes and Steve Johnson. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: Research Genetics web address:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clark, M., Johnson, S.L., Lehrach, H., Lee, R., Li, F., Marra, M., S., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Und, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, Schurk, R., Ritter, Swaller, T., Gibbons, M., Pape, D., Harvey, R., Charles, R., Charl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Putative full length read
The vector to vector length is 97
Seq primer: T3 ET from Amersham.
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Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished Contact: Stephen L. Johnson
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fw20c09.y1 Zebrafish C32 14 somite embryo Danio rerio cDNA clone
IMAGE:5565304 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://www.researchgenetics.com/
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5'-cuacuacuacuaaagcagtggtaacaacgcagagtac-3'. Ends were treated with uracil DNA glycosylase and product with 3 overhangs was annealed to complementary ends of pAMP1. Insert can be excised using EcoRI and NotI. Library constructed by Joe Barnes and Steve Johnson (Washington)
                                                                                                                                                                                                   /clone lib="Zebrafish C32 14 somite embryo" /note="Yector: pAMP1; Site 1: EcoRI; Site 2: Not strand cDNA synthesis was primed using oligo-dT magnetic beads with an additional primer 5'-gcggccgctaatacgactcacta-taggg-3'. Second stransynthesis was a 3-cycle PCR using the primers
                                                                                                                                                                     5'-caucaucaucauggccgctaatacgactcactataggg-3' and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="embryo, 14 somite"
/lab_host="DH10B"
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Pred. No. 7.9e+04;
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          Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                      BZ352958 100 bp DNA linear GSS SALK 119560.52.85.x Arabidopsis thaliana TDNA insertion Arabidopsis thaliana genomic clone SALK_119560.52.85.x,
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                                                       Arabidopsis thaliana
                                                                                     GSS.
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BZ352958
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                                                                Arabidopsis thaliana (thale cress)
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BP 191 91006 EVRY cedex - FRANCE
- Web : www.genoscope.cns.fr)
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Arabidopsis thaliana genome survey sequence T7 end of BAC T6P22 of
TAMU library from strain Columbia of Arabidopsis thaliana, genomic
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/mol_type="genomic DNA'
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A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished Contact: Joseph R. Ecker Salk Institute Genomic Analysis Laboratory (SIGnAL) Salk Institute for Biological Studies The Salk Institute for Biological Studies 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA Tel: 858 453 4100 x1752 Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This is single pass sequence recovered from the left border of TDNA. This sequence lies within an annotated intron of At3g27020. Class: TDNA tagged.

Location/Qualifiers
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                                                                                                                                                                                                                                                                                                      /organism="Arabidopsis thaliana"
/organism="Golumbia 0"
/strain="Columbia 0"
/db_xref="taxon:3702"
/clone="SALK_119560.52.85.x"
/clone="SALK_119560.52.85.x"
/clone="PCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
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